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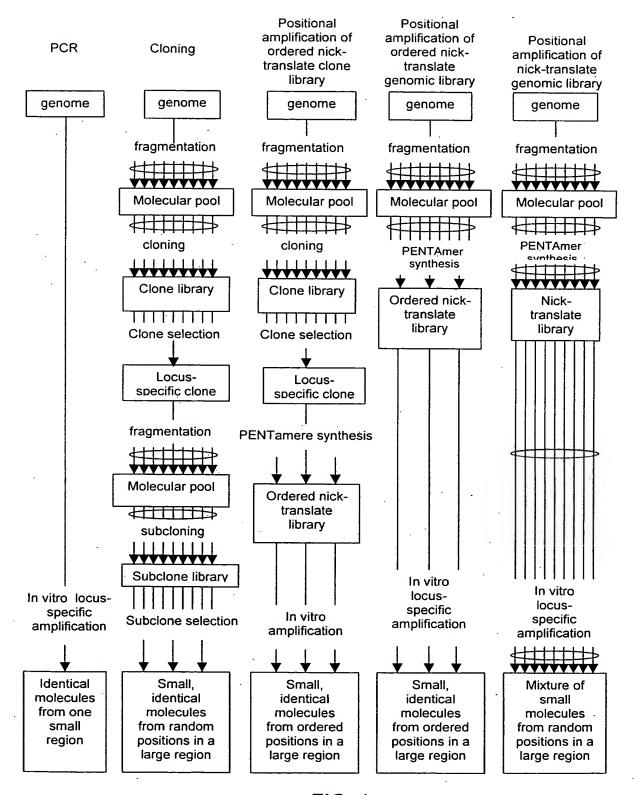


FIG. 1

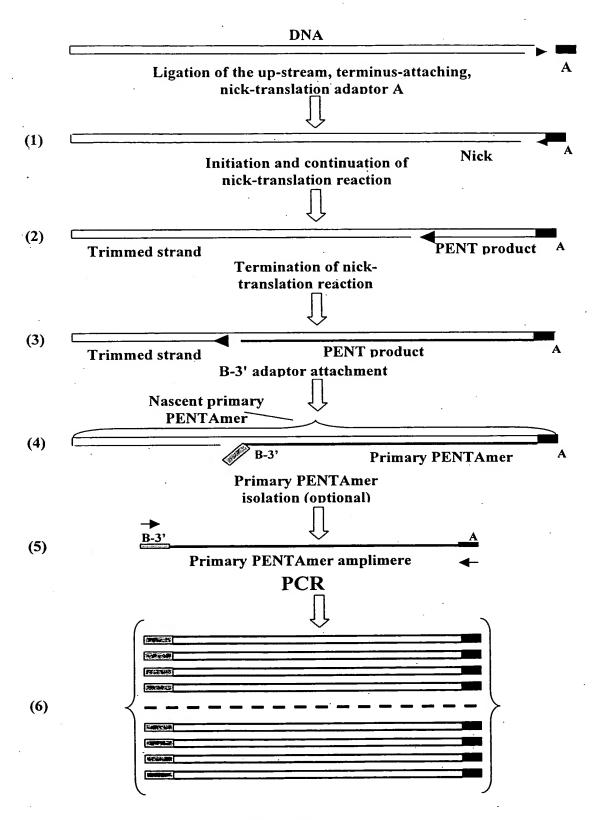


FIG. 2A

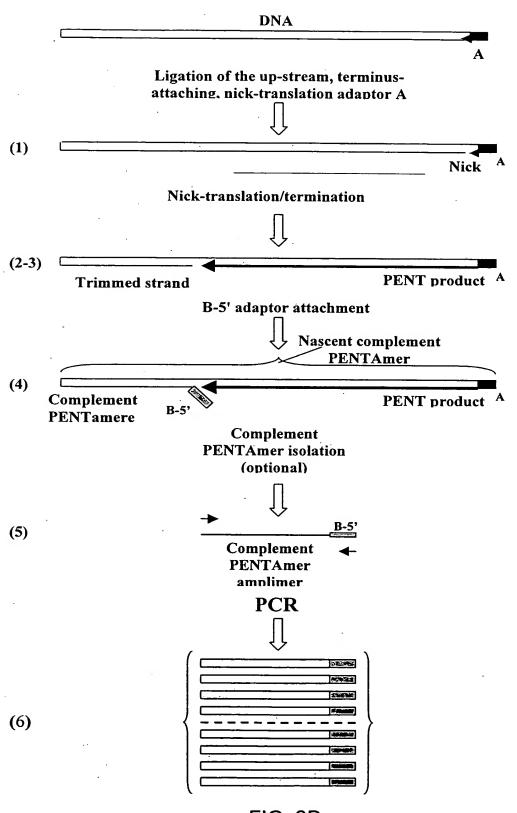


FIG. 2B.

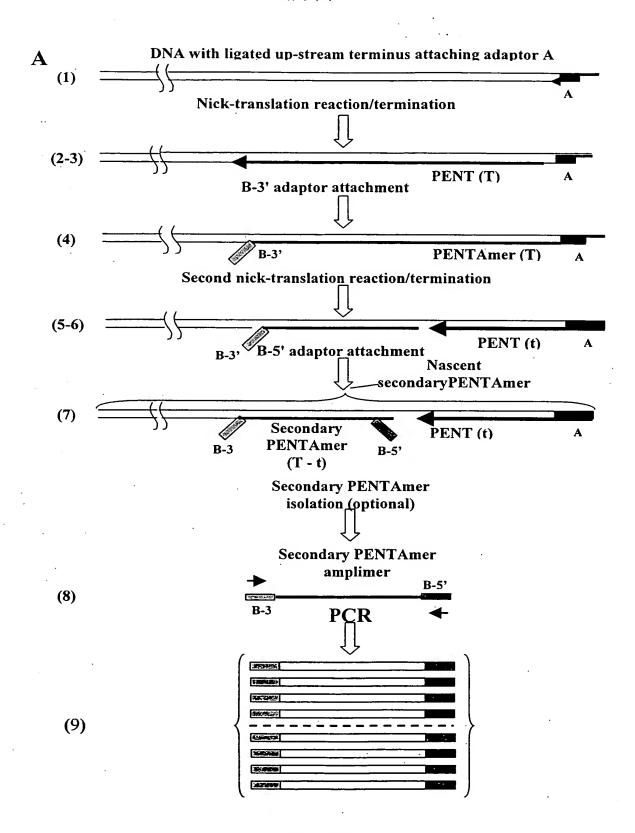


FIG. 3A

B (1-3) as in FIG. 3A

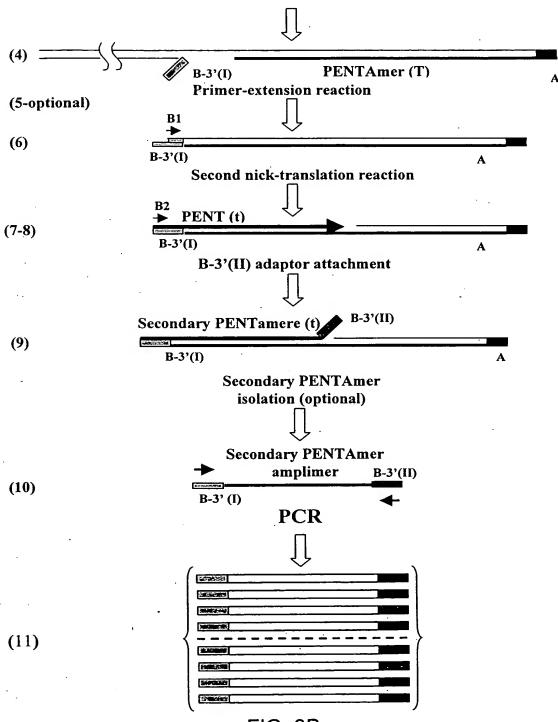


FIG. 3B

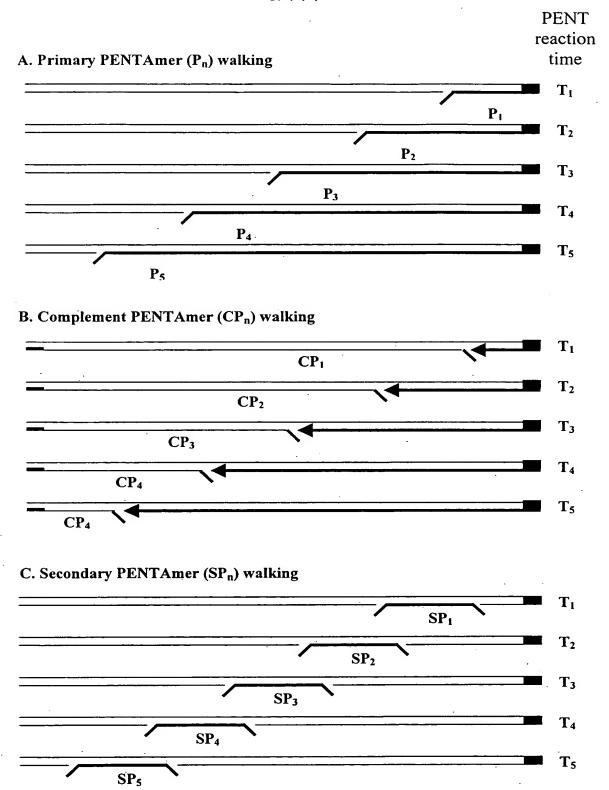


FIG: 4

FIG. 5

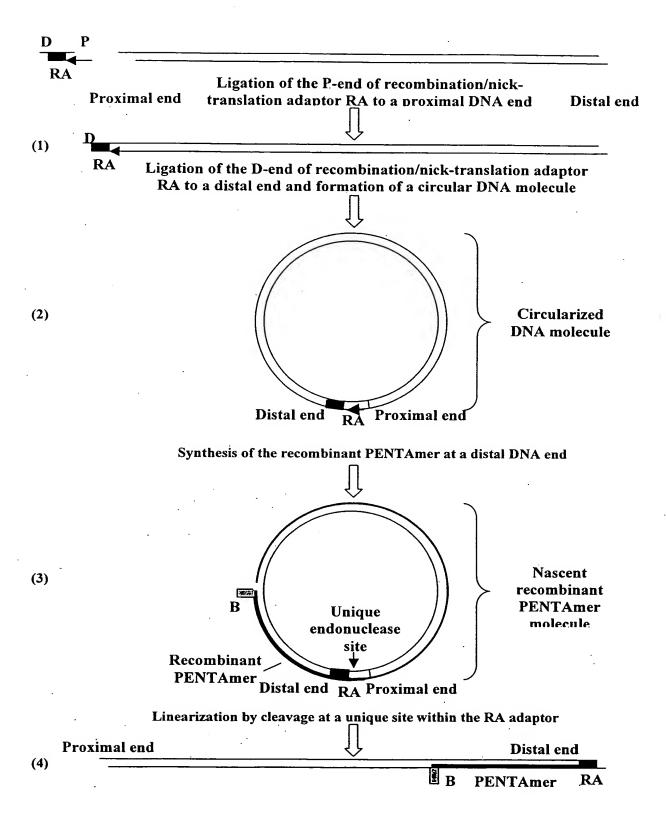


FIG. 6

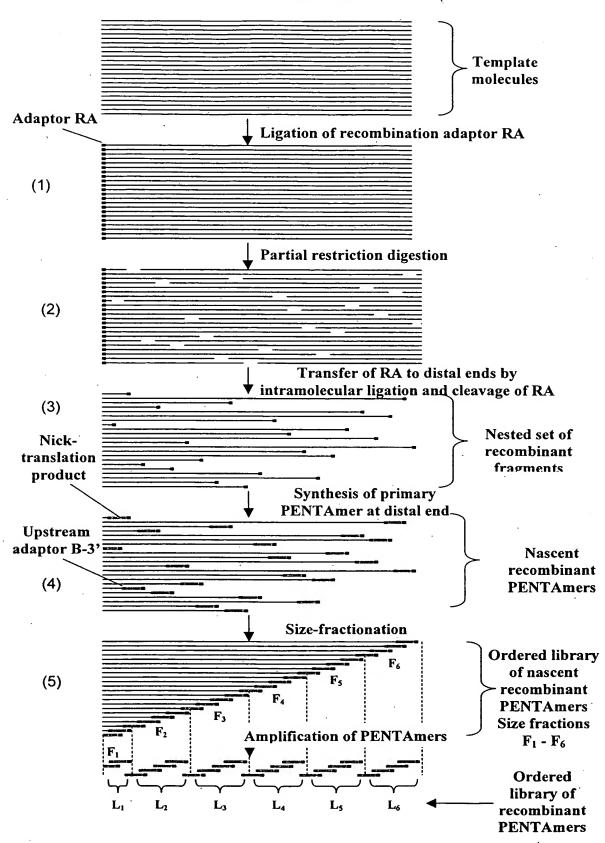


FIG. 7

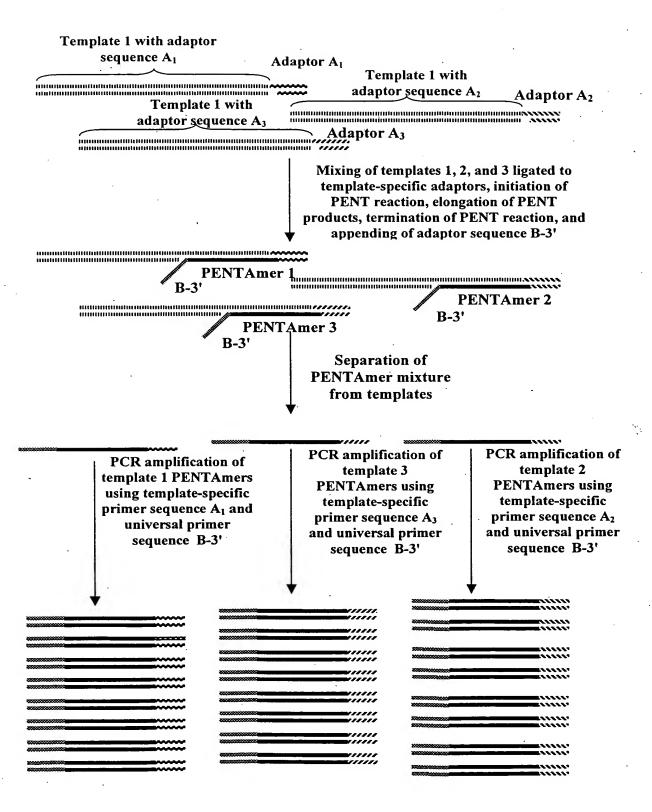
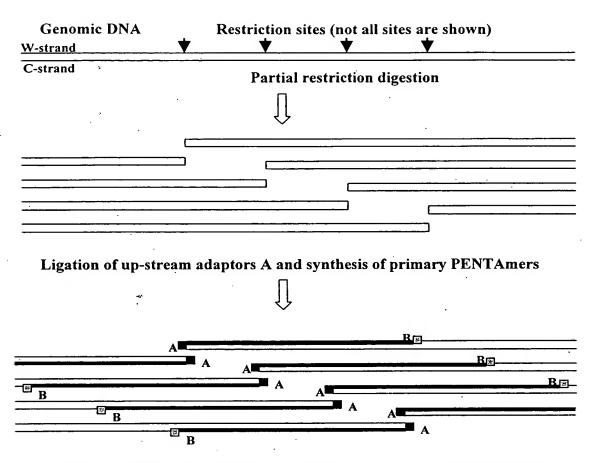


FIG. 8



Primary PENTamere isolation by affinity capture or by alkaline gel size

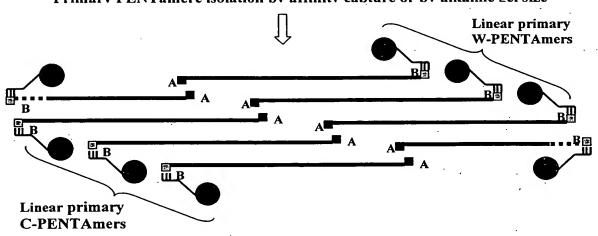


FIG. 9A

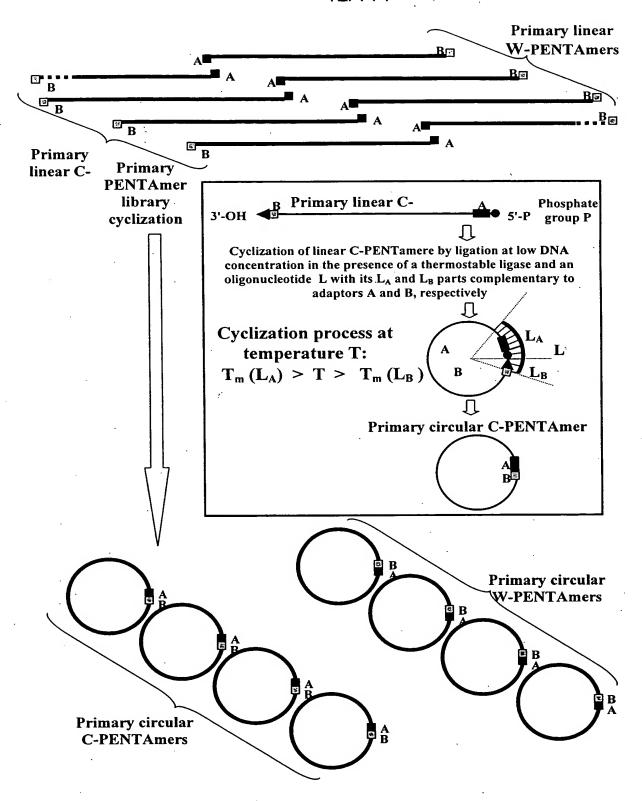


FIG. 9B

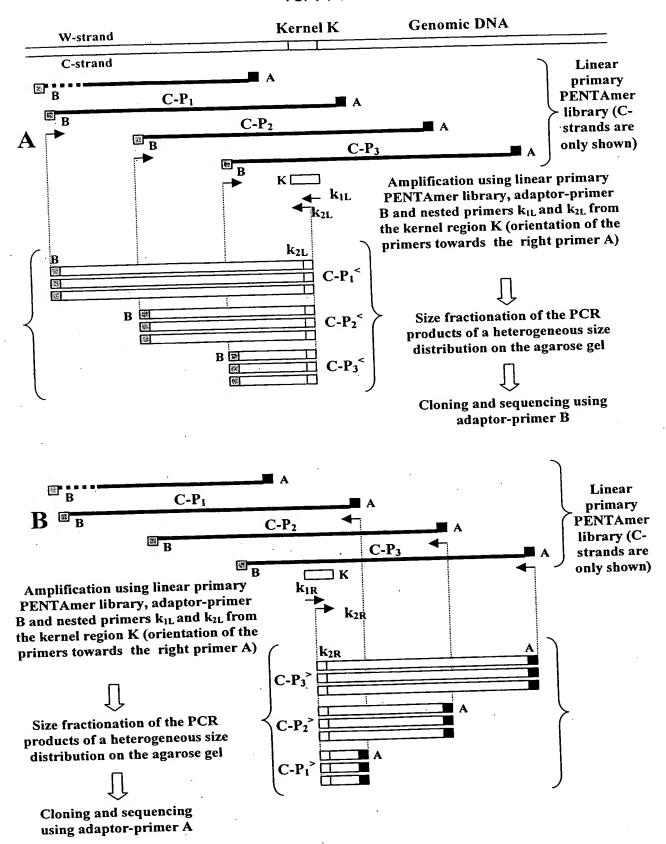


FIG. 10

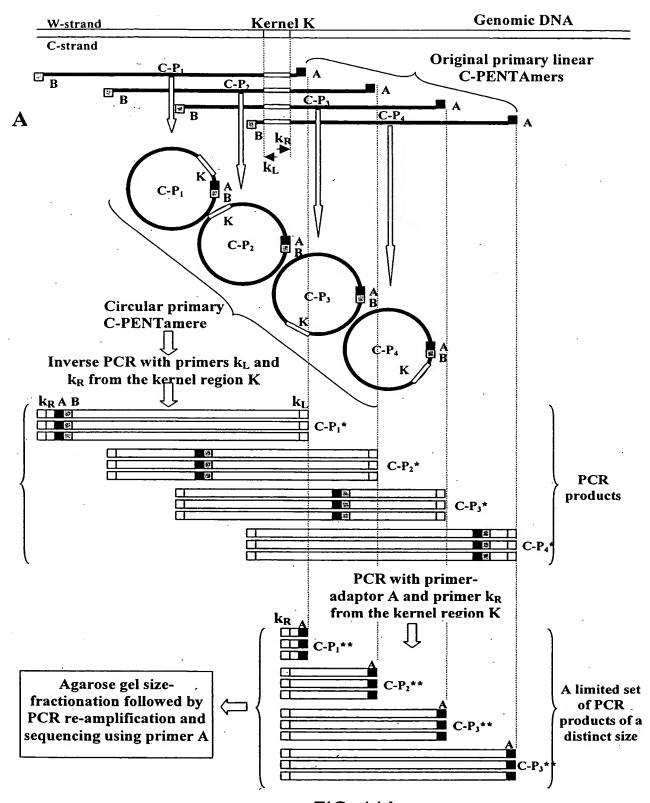


FIG. 11A

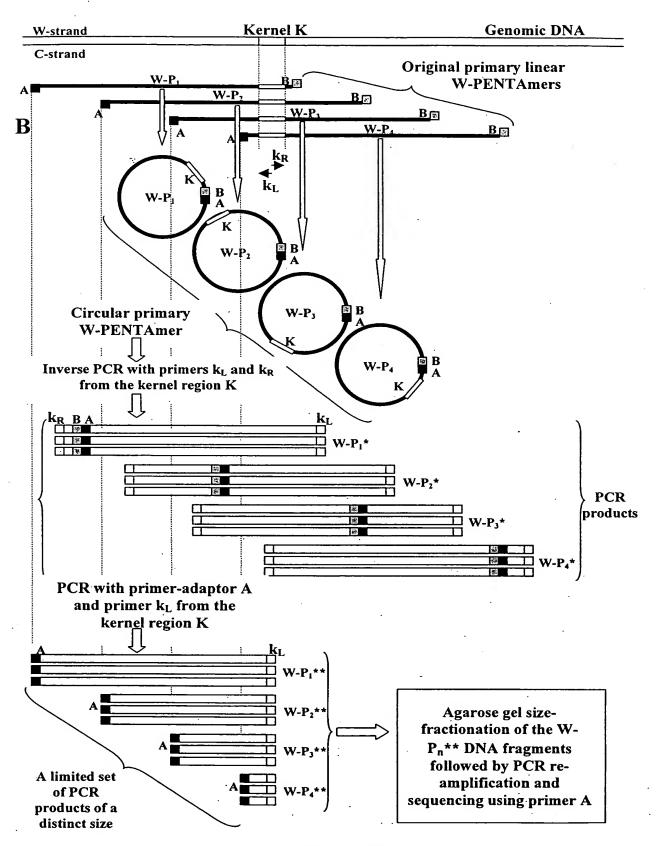


FIG. 11B

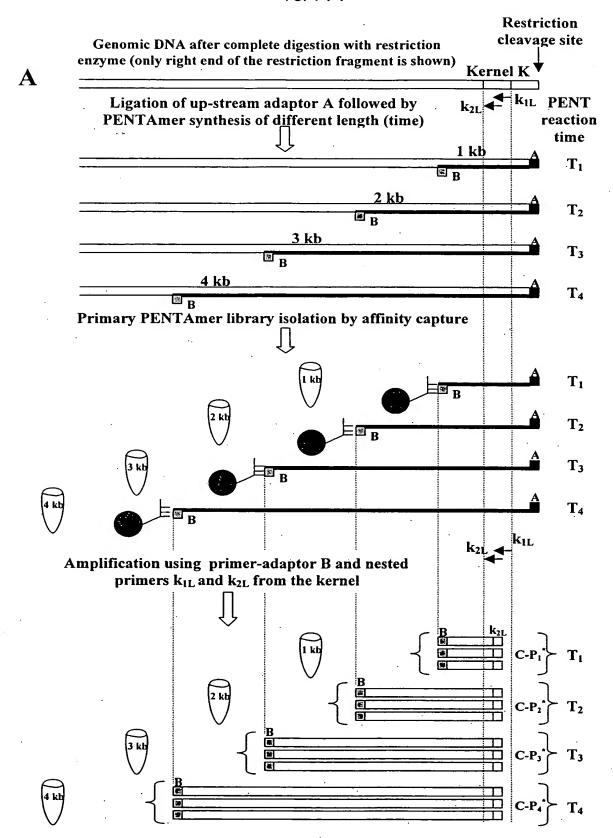
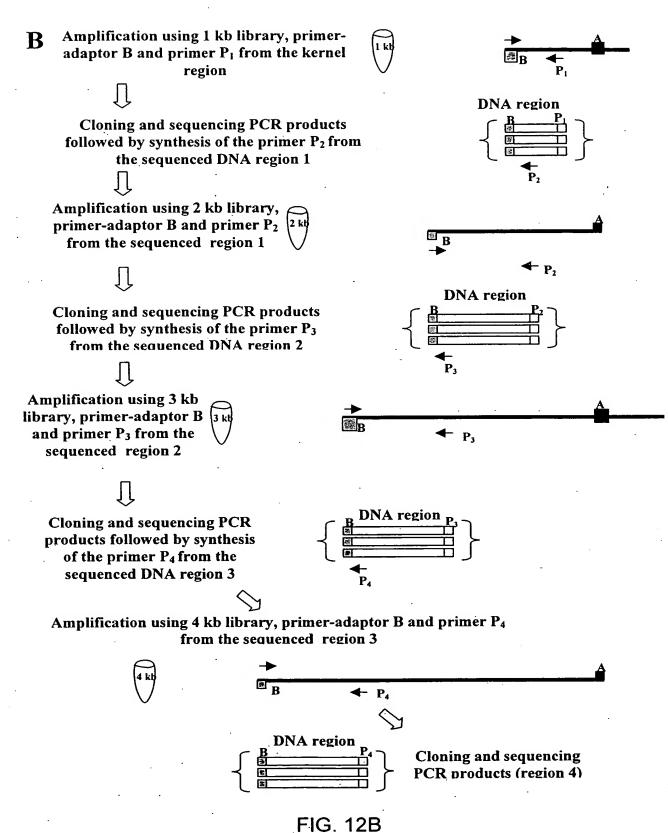


FIG. 12A



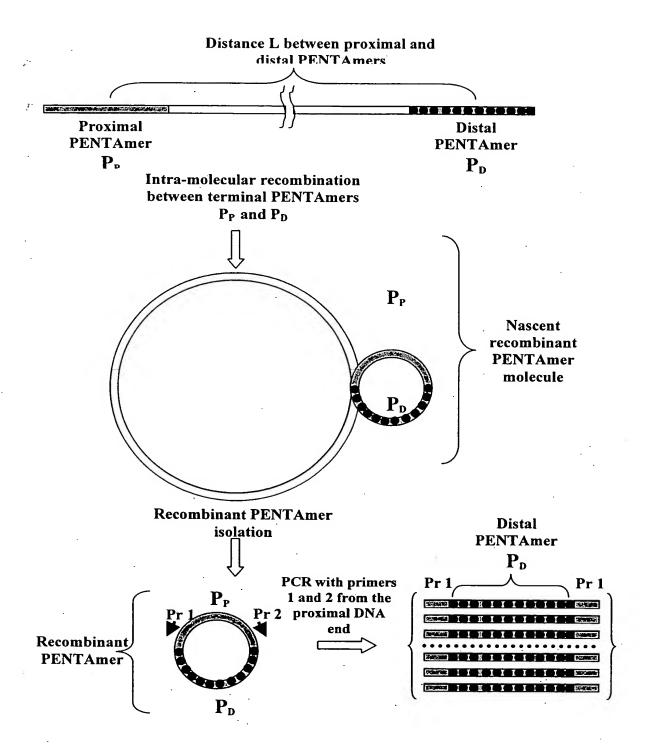
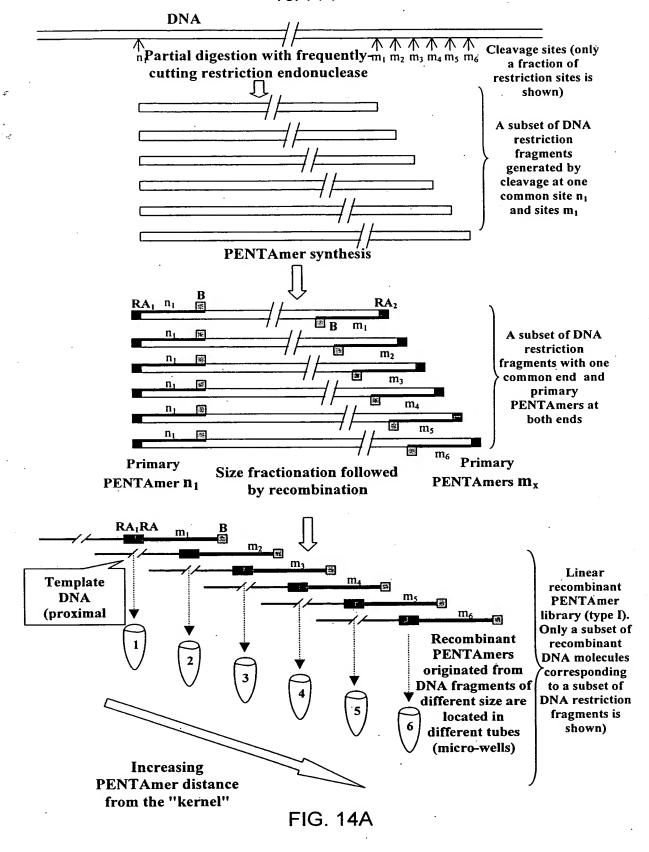


FIG. 13



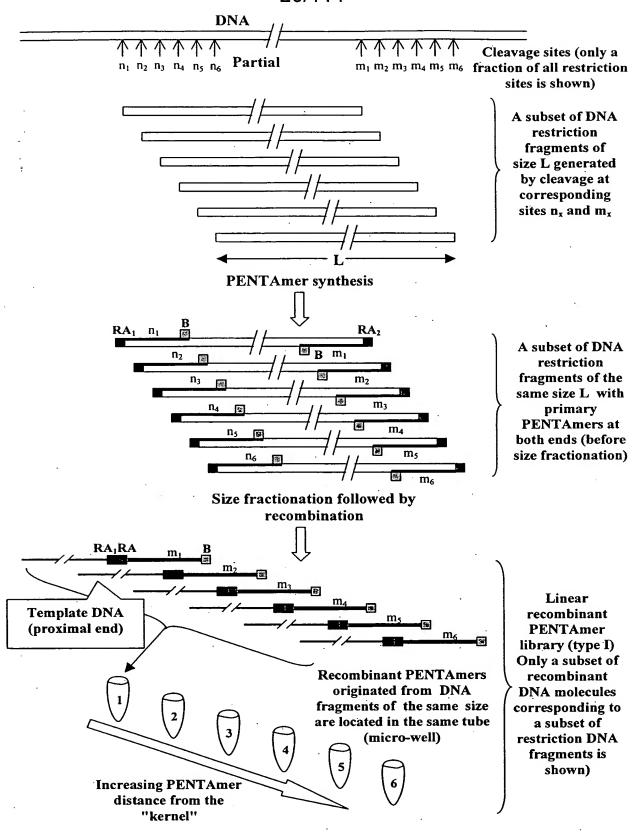
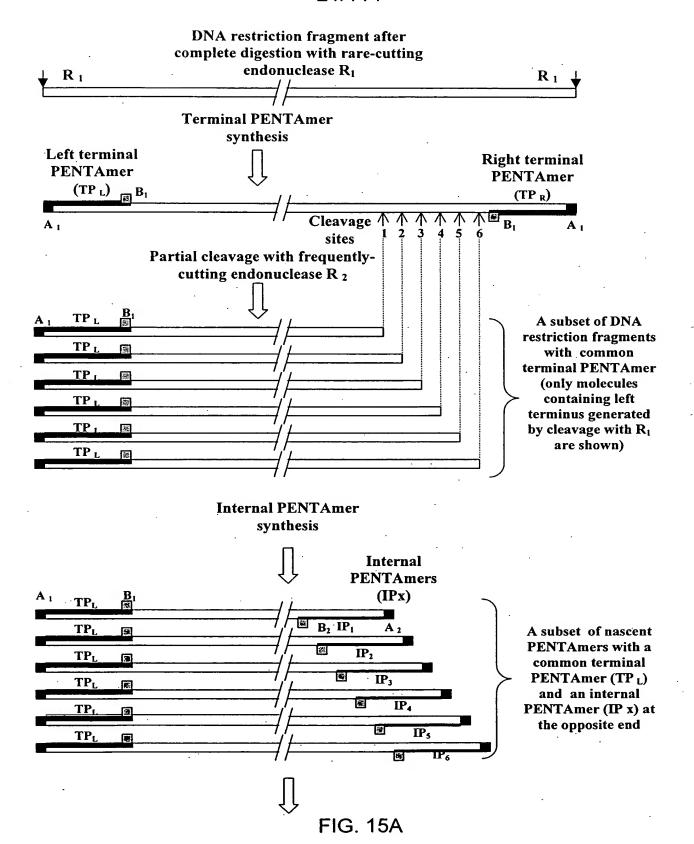
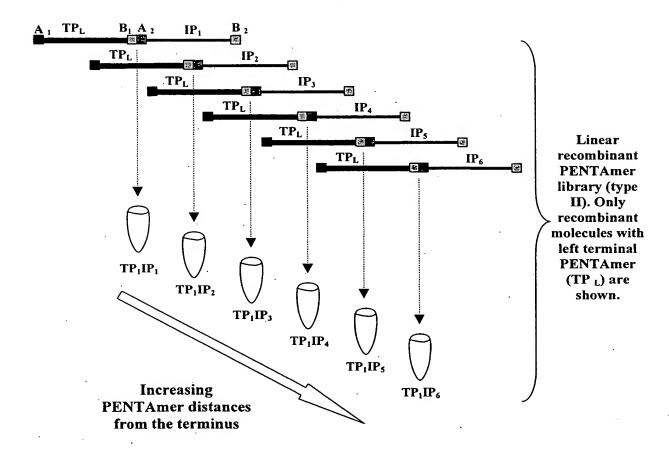


FIG. 14B



Size fractionation followed by recombination





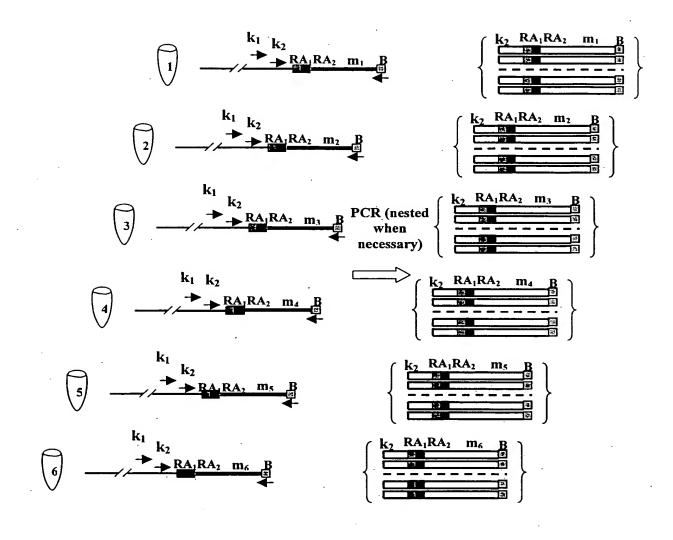
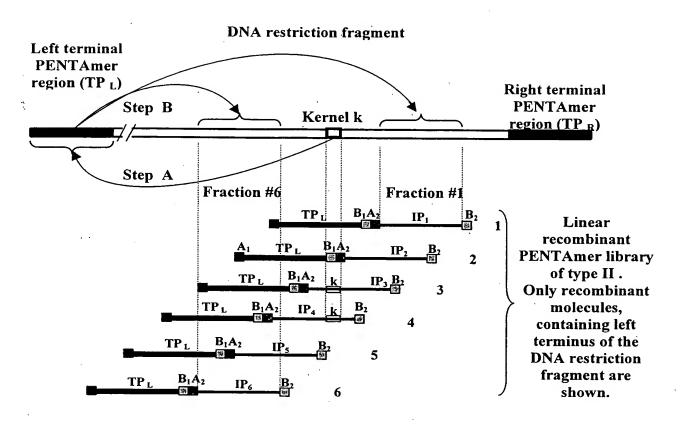


FIG. 16



Step A. Amplification of terminal PENTAmers (TP) of a DNA restriction fragment using "One-tube" unordered library II, nested primers K₁ and K₂ from the internal "kernel" region k and adaptor-primer A₁ (only amplification of left terminal PENTAmer is shown)

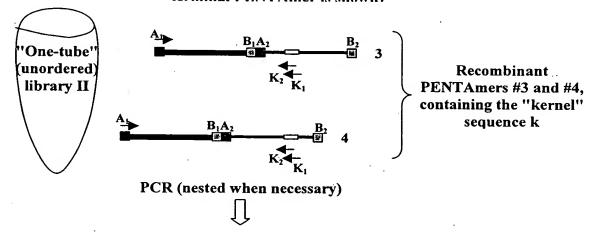
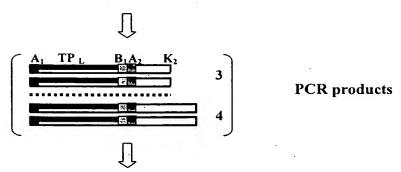


FIG. 17A



Sequencing left terminal PENTAmer DNA (TP $_L$) using primers A_1 and/or B_1 followed by selection and construction of the PCR primers T_1 and T_2 from the terminal PENTAmer region



Step B. Amplification of the internal PENTAmer (IP) fractions of the DNA restriction fragment using linear ordered library II, nested primers T₁ and T₂ from the terminal PENTamere region TP and the adaptor-primer B₂

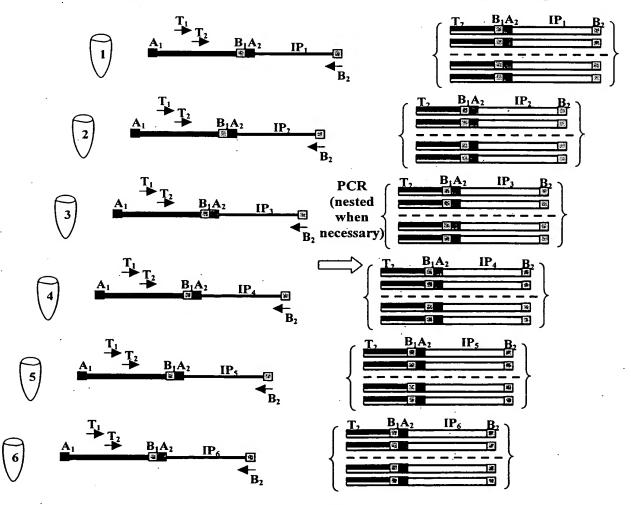
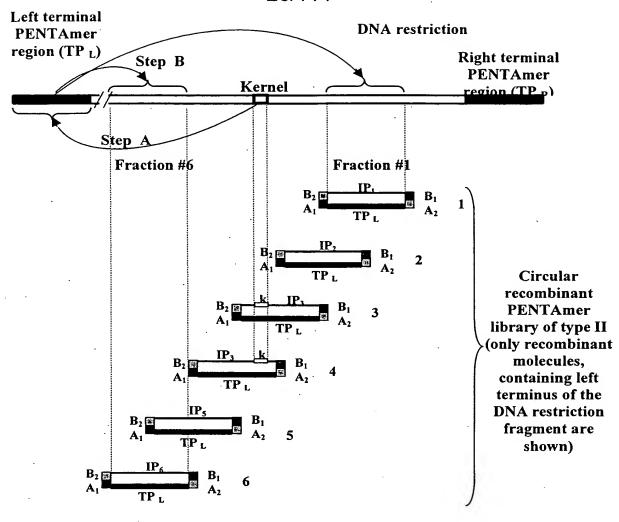


FIG. 17B



Step A. Amplification of terminal PENTAmers (TP) of a DNA restriction fragment using "One-tube" circular library II, inverse primers K_1 and K_2 from the internal "kernel" region k

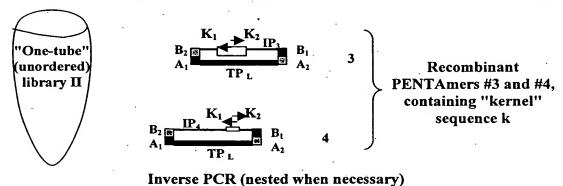
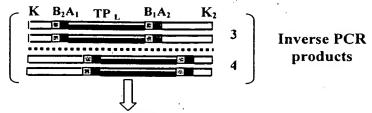


FIG. 17C



Sequencing a terminal PENTAmer DNA (TP) using primers A_1 and /or B_1 followed by selection and construction of two inverse PCR primers T_1 and T_2

Step B. Amplification of the internal PENTAmer (IP) fractions of the DNA restriction fragment using inverse primers T₁ and T₂ from the terminal PENTAmer region and ordered circular library II

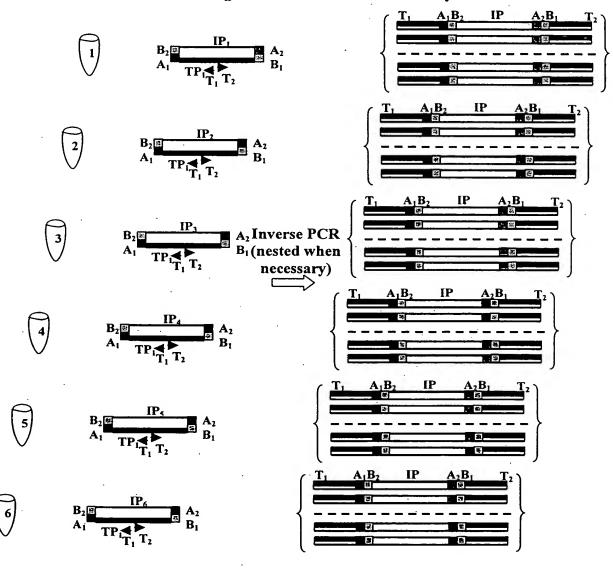
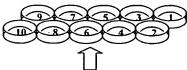


FIG. 17D

A. SmartGenome DNA library I



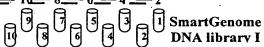
"Positional" cloning and sequencing ordered DNA fragments located right next to the kernel



PCR-amplified

Left of the kernel

DNA fragment = 10 = 8 nools



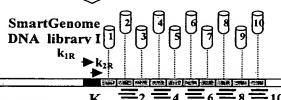


Amplification of ordered DNA fragment pools located to the left of the kernel region K, using SmartGenome DNA library I and "left-oriented" primers k1L and k2L from the K-region.



Amplification of ordered DNA fragment pools located to the right of the kernel region K, using SmartGenome DNA library I and 'right-oriented " primers k1R and k2R from the K-region.



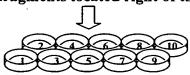


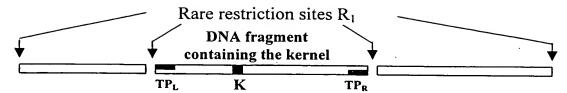
Right of the kernel

1三3三

PCR-amplified DNA fragment pools

"Positional" cloning and sequencing ordered DNA fragments located right of the kernel

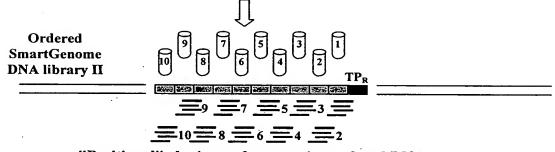




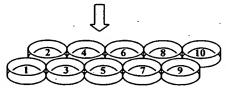
Amplification, isolation and sequencing of termini TP_L and TP_R of the DNA restriction fragment R with the kernel sequence using unordered "One-tube" SmartGenome DNA library II and primers k_1 and k_2 from the K-region



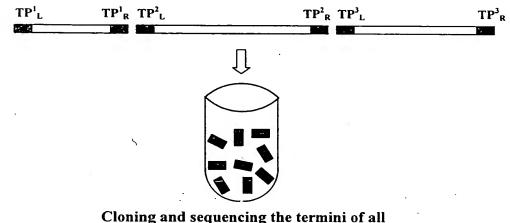
Amplification of the ordered internal DNA molecules located within the same DNA restriction fragment R using ordered SmartGenome DNA library II and primers from the terminal regions



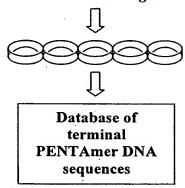
"Positional" cloning and sequencing ordered DNA molecules within the restriction fragment containing the kernel



Step 1 Linear amplification of termini (TP_L and TP_R) of all DNA restriction fragments using unordered "One-tube" SmartGenome DNA library II and adaptor-primers A₁ and B₁ (see Fig. 17A)



Cloning and sequencing the termini of all DNA restriction fragments



Step 2 Amplification of the ordered internal DNA fractions located within the DNA restriction fragments using ordered SmartGenome DNA library II and primers from the terminal regions

Step 3 Amplification, isolation and sequencing of the "linking" DNA fractions using ordered SmartGenome DNA library I and primers within the adjacent terminus region

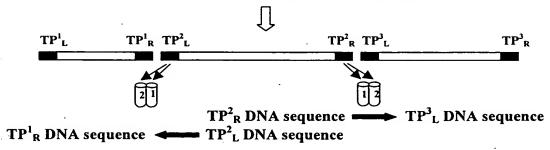


FIG. 18C

31/114

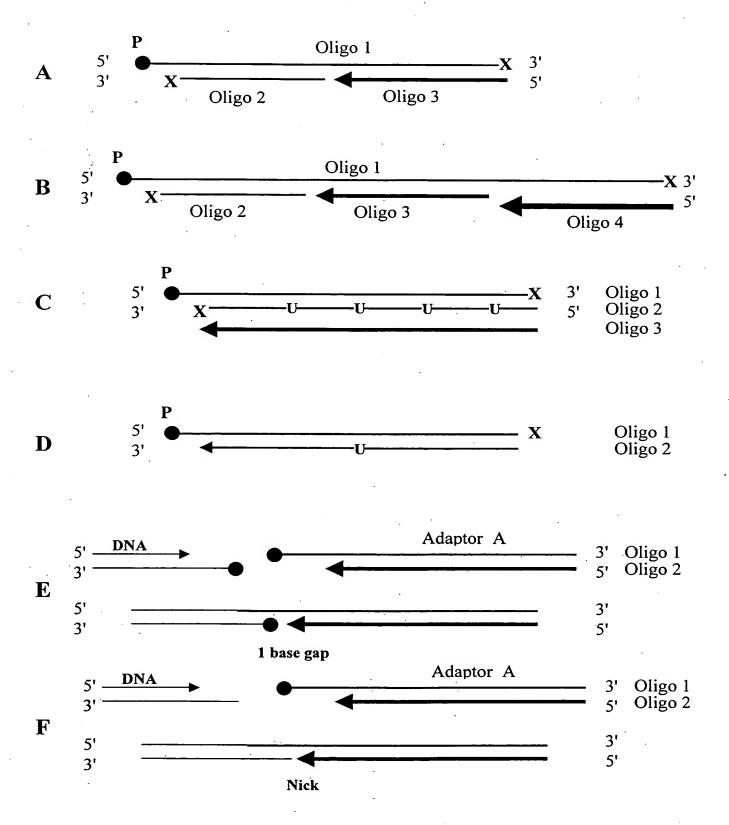


FIG. 19

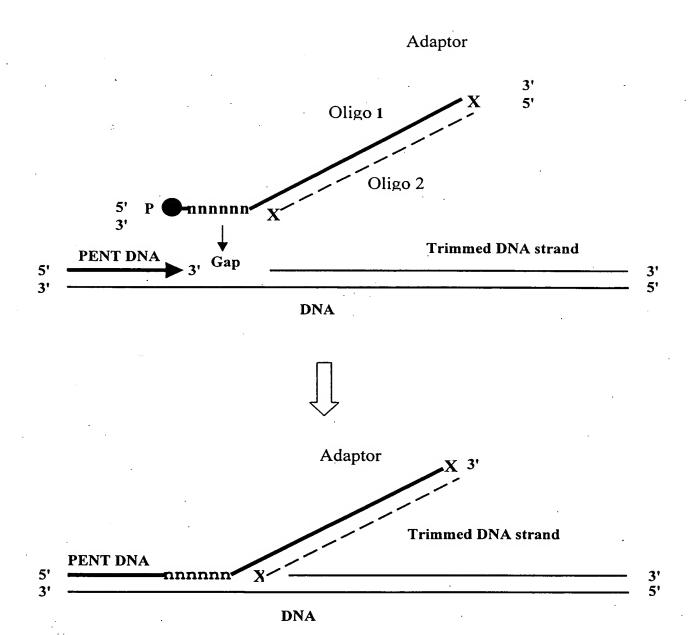


FIG. 20

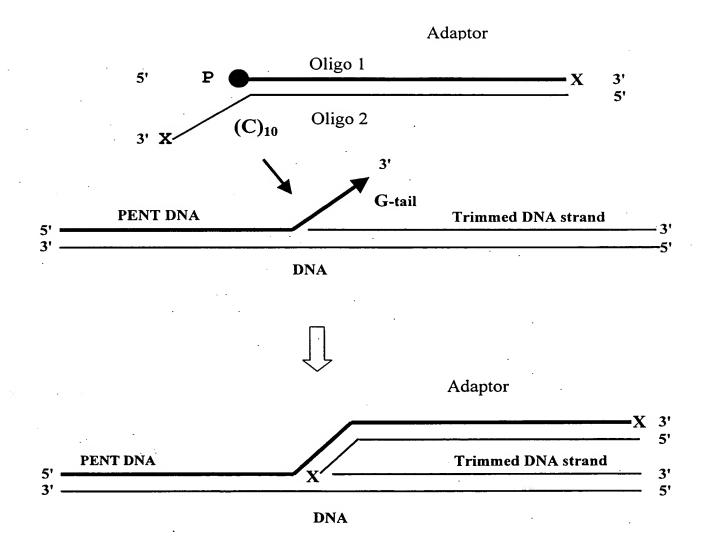


FIG.21

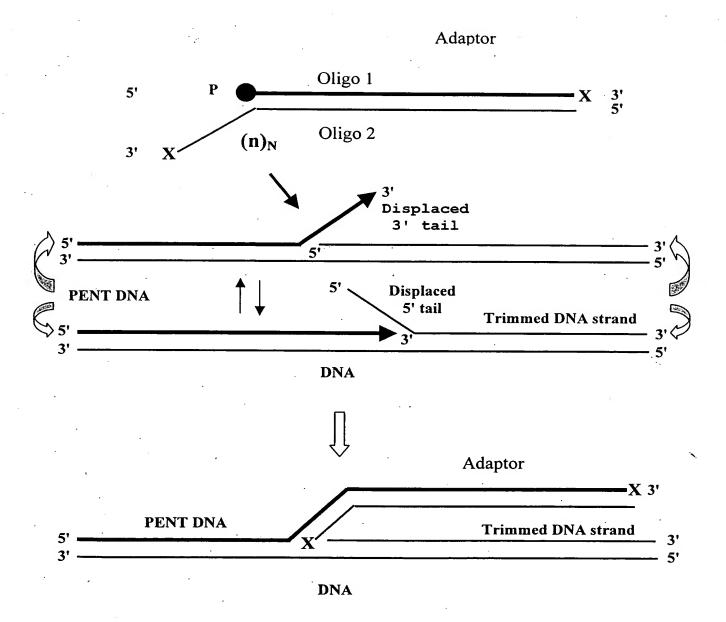


FIG. 22

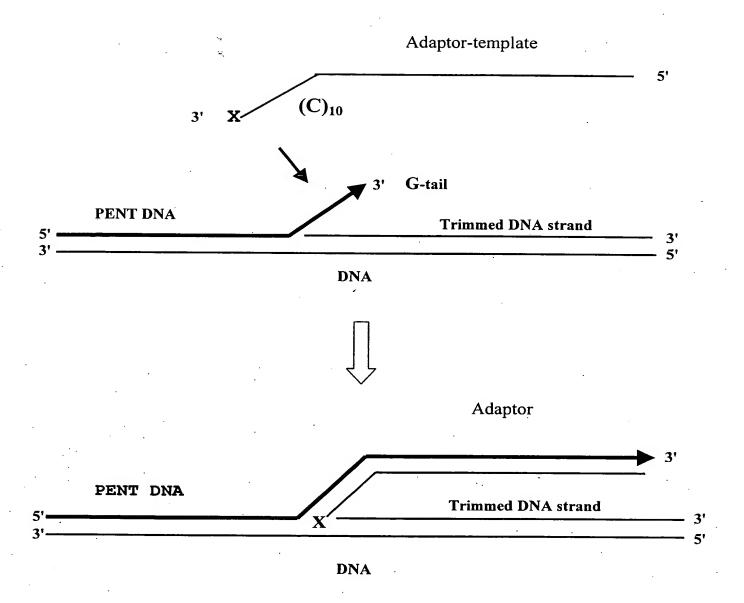


FIG. 23

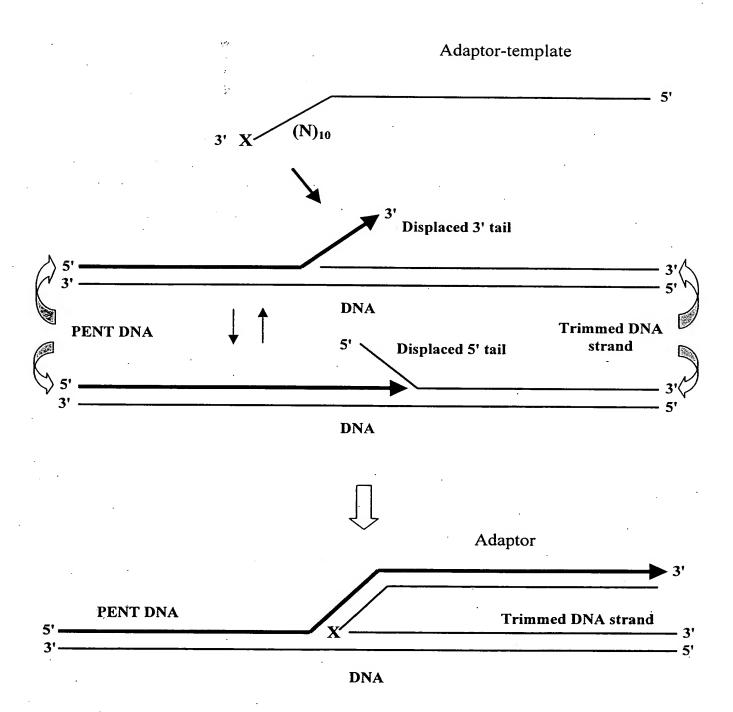


FIG. 24

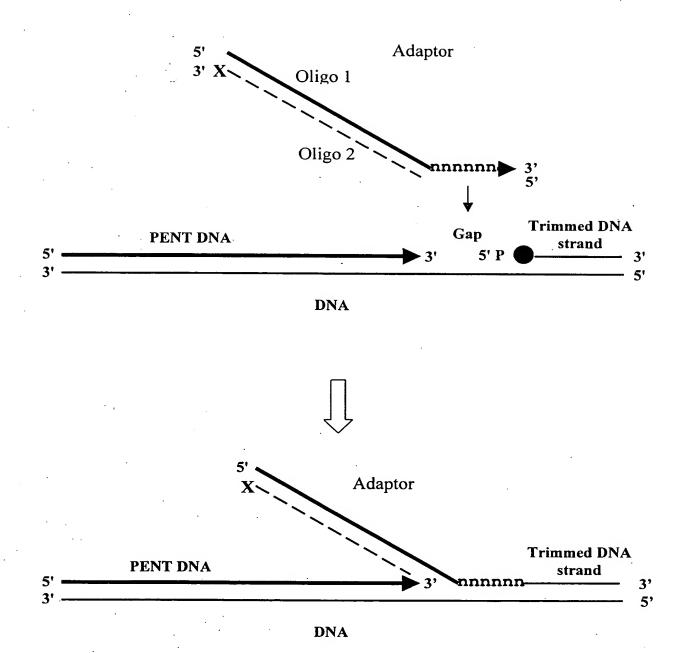


FIG. 25

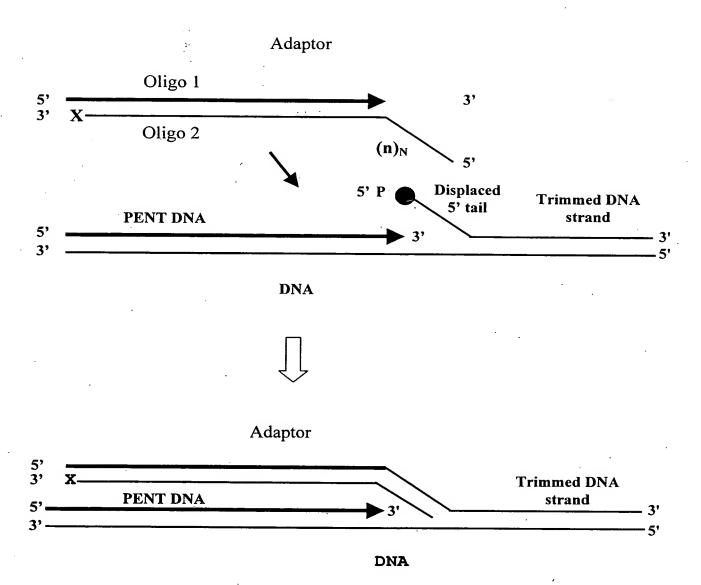
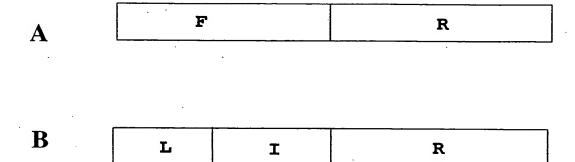
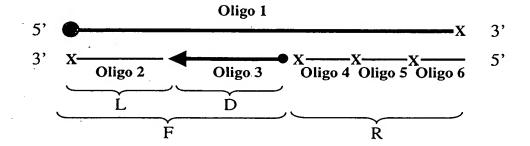


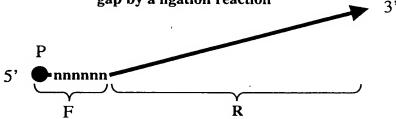
FIG. 26



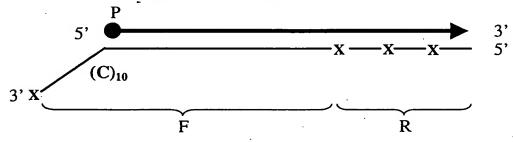
A Up-stream terminus-attaching nick-translation recombination adaptor RA



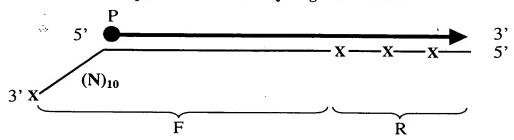
B Down-stream nick-attaching recombination adaptor RB-3' (I) targeted to a gap by a ligation reaction



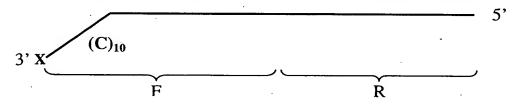
C Down-stream nick-attaching recombination adaptor RB-3' (II) targeted to a poly-G tail by a ligation reaction



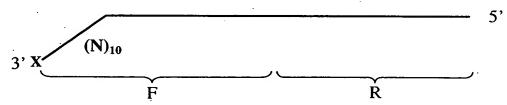
D Down-stream nick-attaching recombination adaptor RB-3' (III) targeted to a displaced 3' DNA tail by a ligation reaction



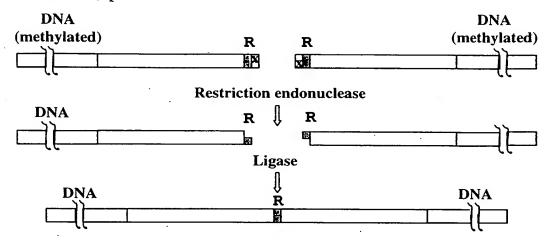
E Down-stream nick-attaching recombination adaptor RB-3' (IV) targeted to a poly-G tail as a template for a polymerization-extension reaction



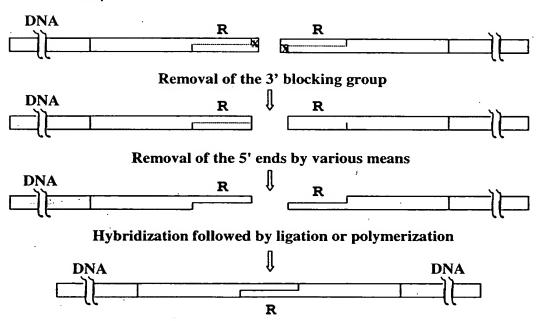
F Down-stream nick-attaching recombination adaptor RB-3' (V) targeted to a displaced 3' DNA tail as a template for a polymerization-extension reaction

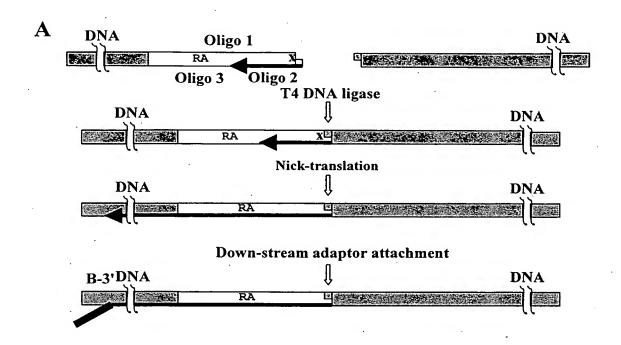


A RecAdaptors-Class I



B RecAdaptors-Class II





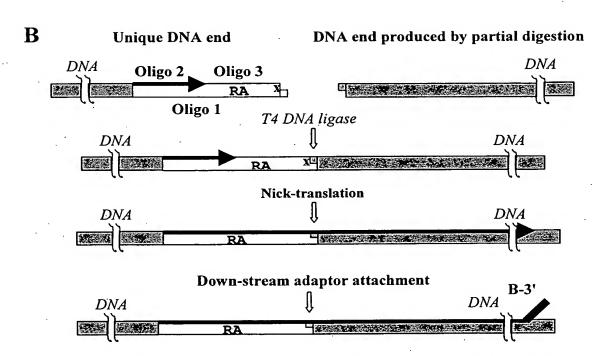


FIG. 30B

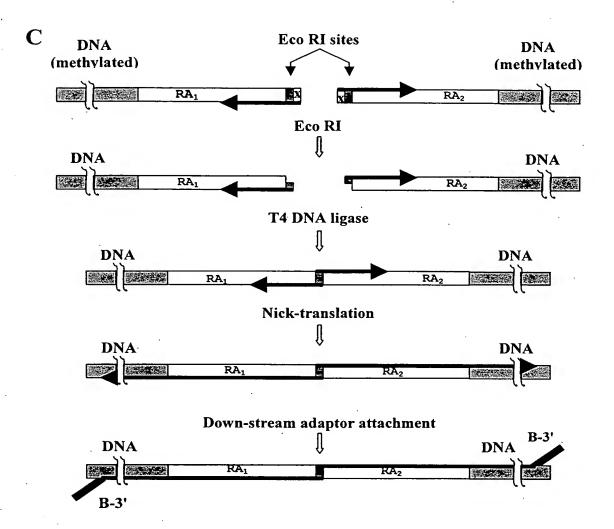


FIG. 30C

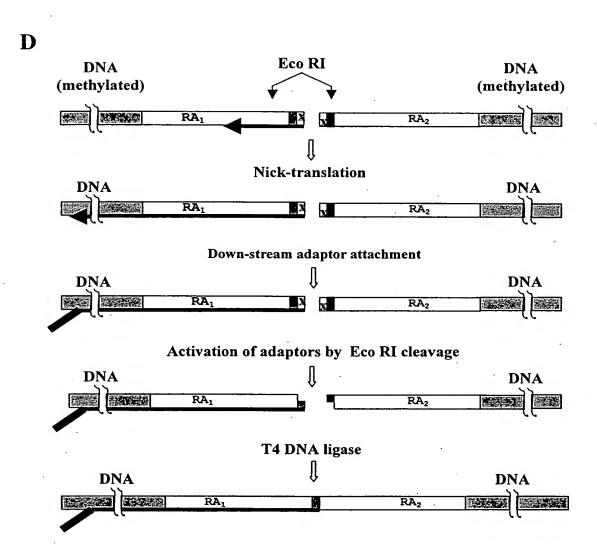


FIG. 30D

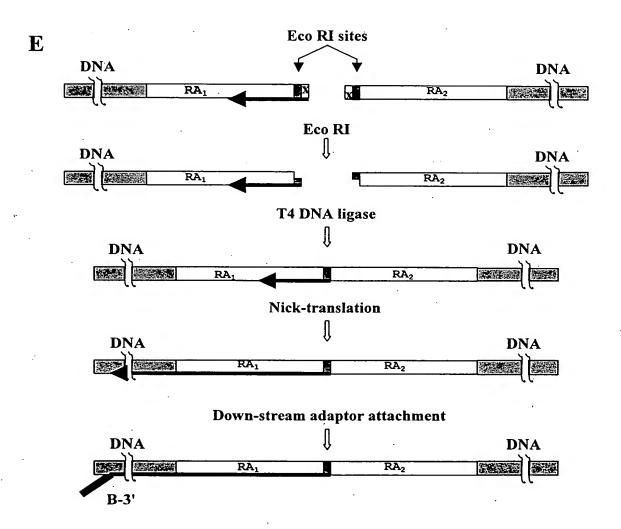


FIG. 30E

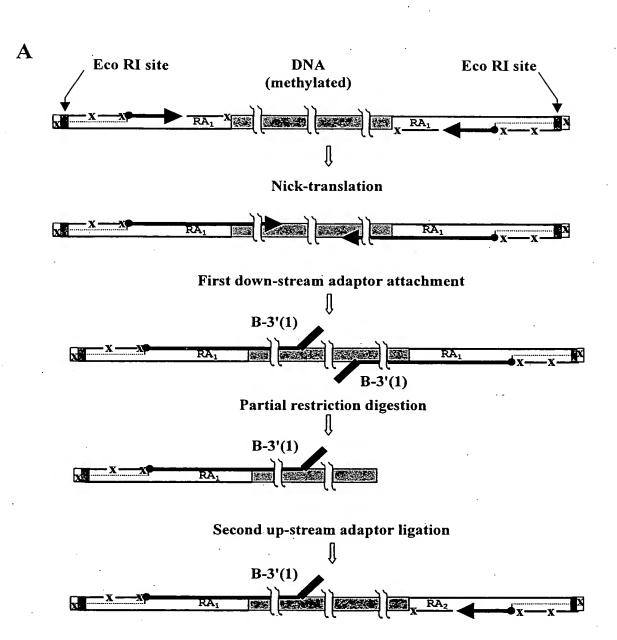


FIG. 31A

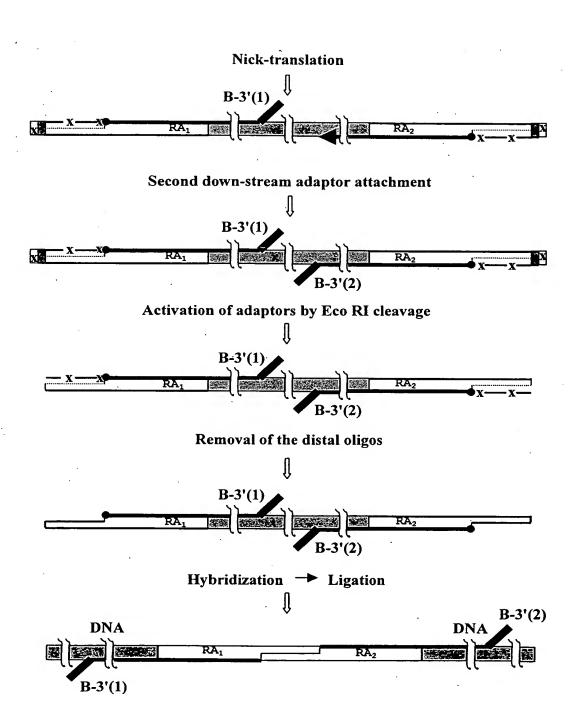


FIG. 31B

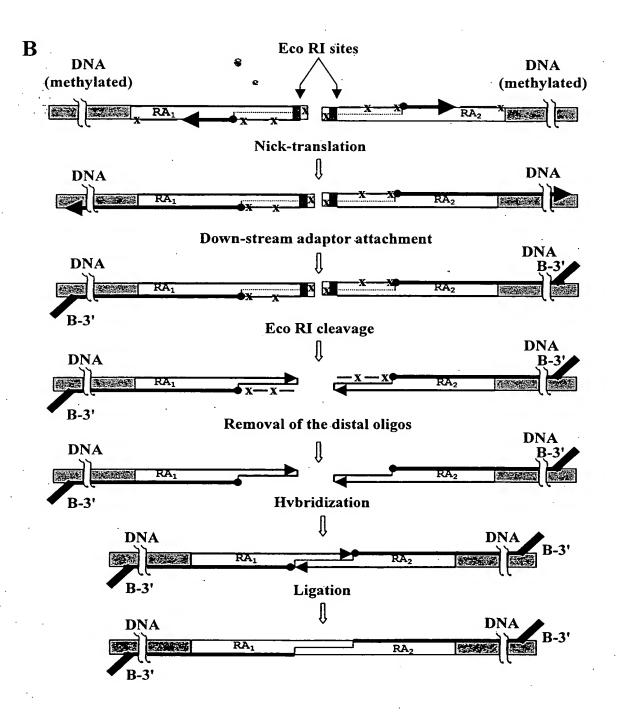


FIG. 31C

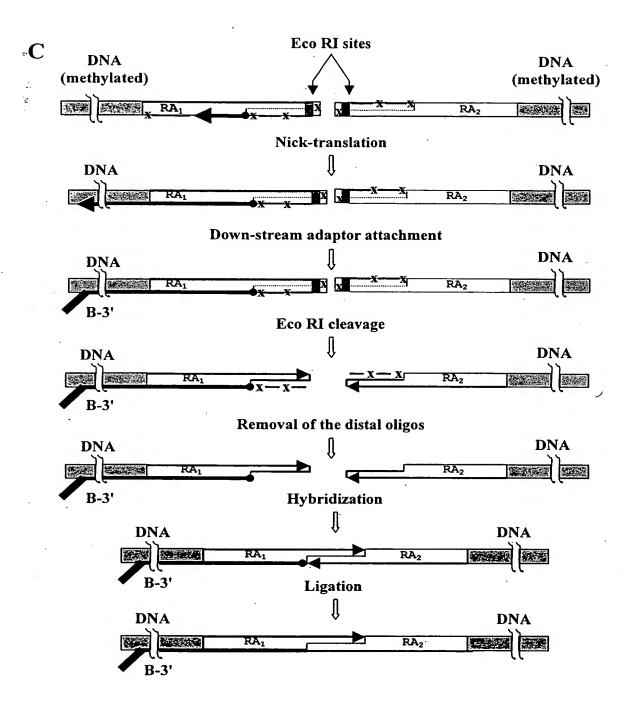


FIG. 31D

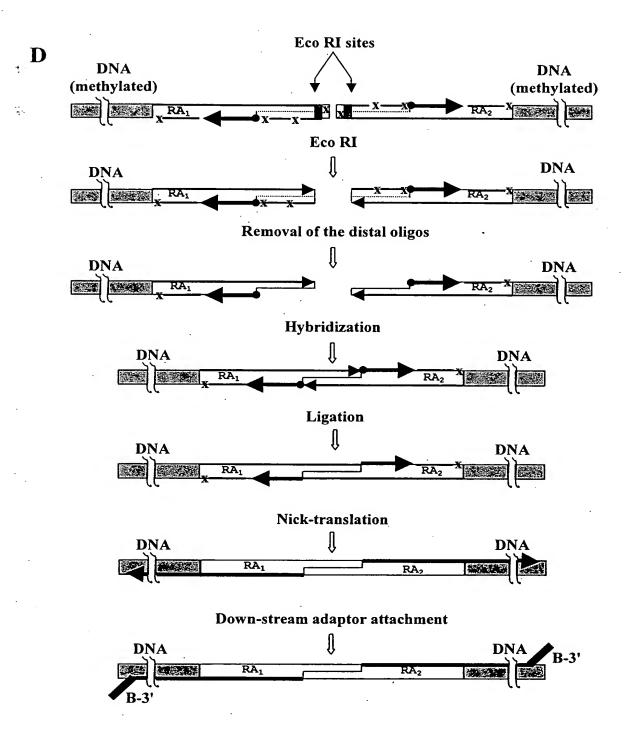


FIG. 31E

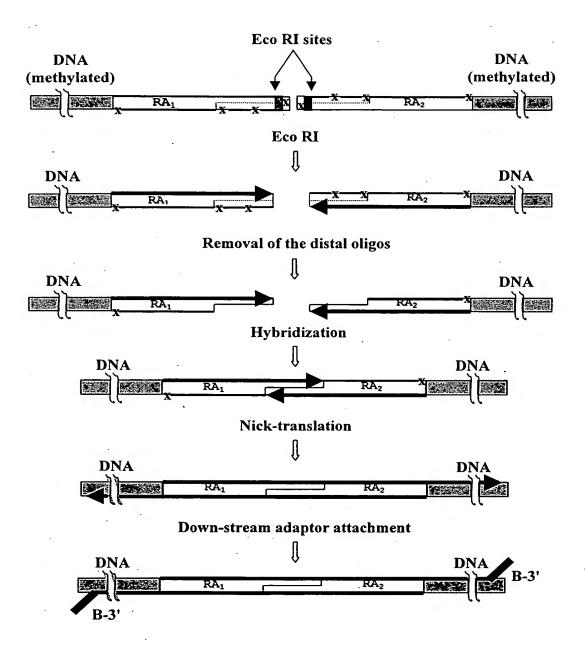


FIG. 32

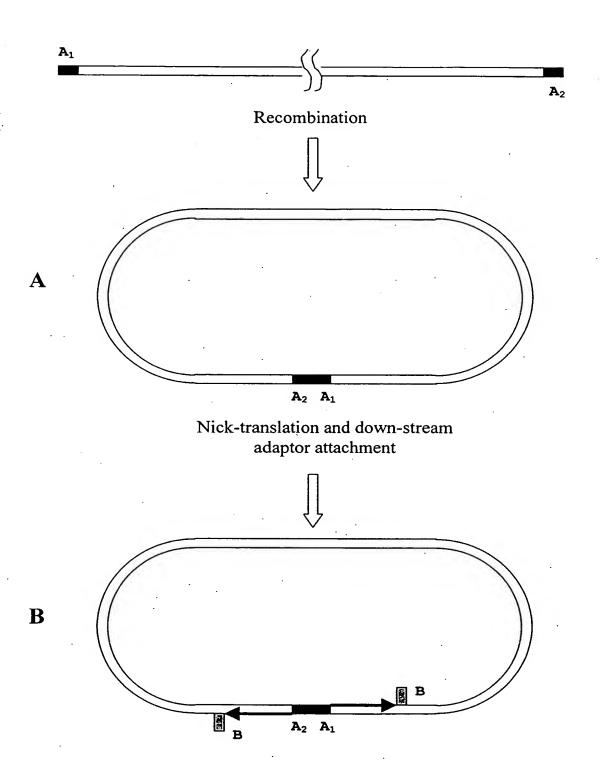


FIG. 33

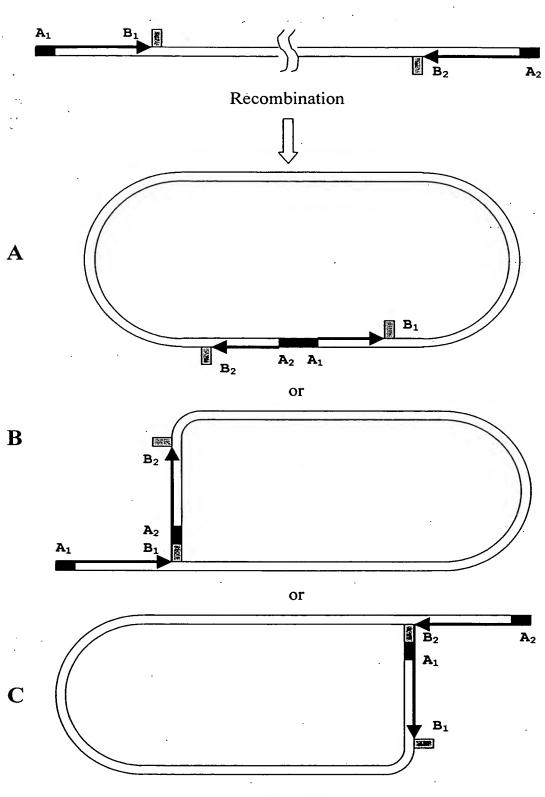
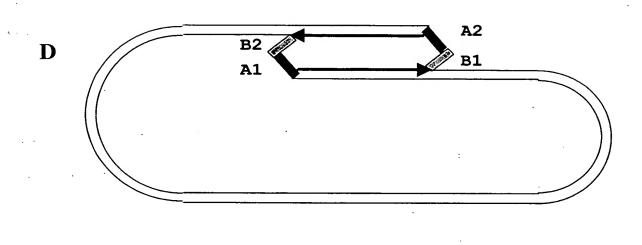
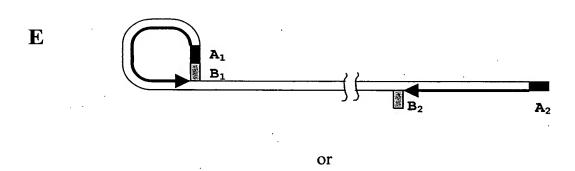


FIG. 34A

or



or



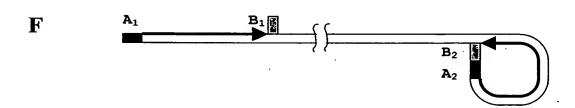


FIG. 34B

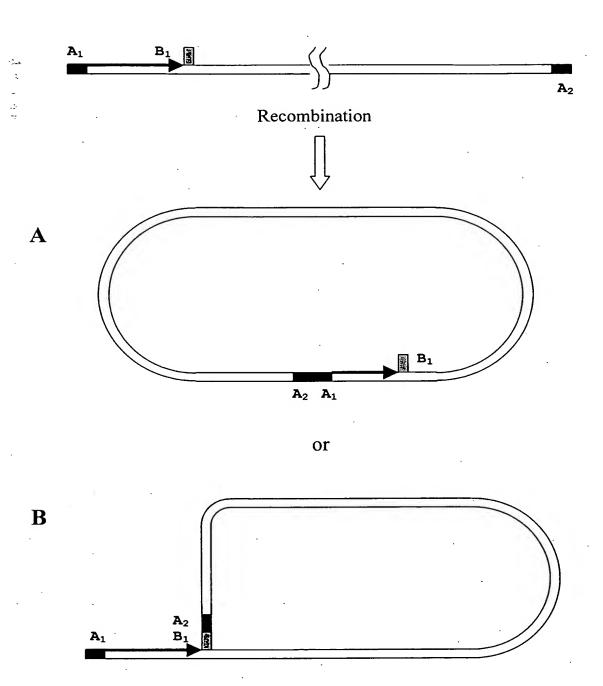


FIG. 35

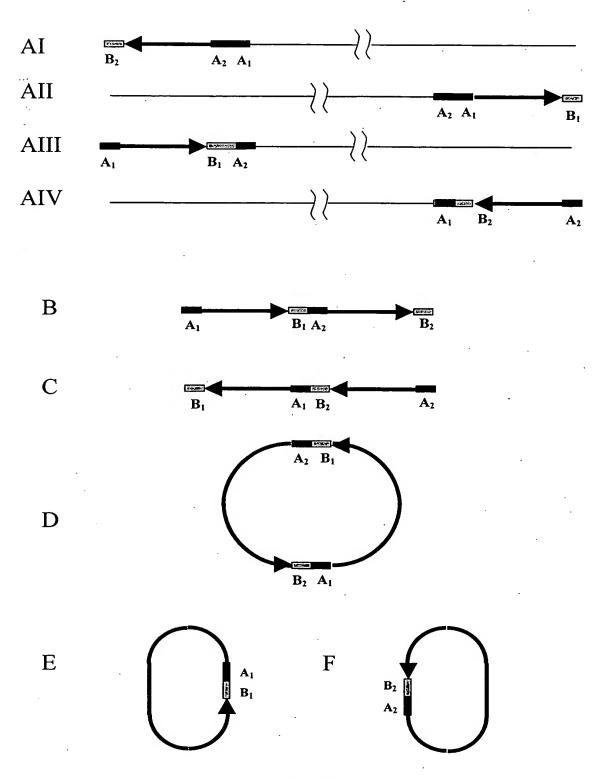
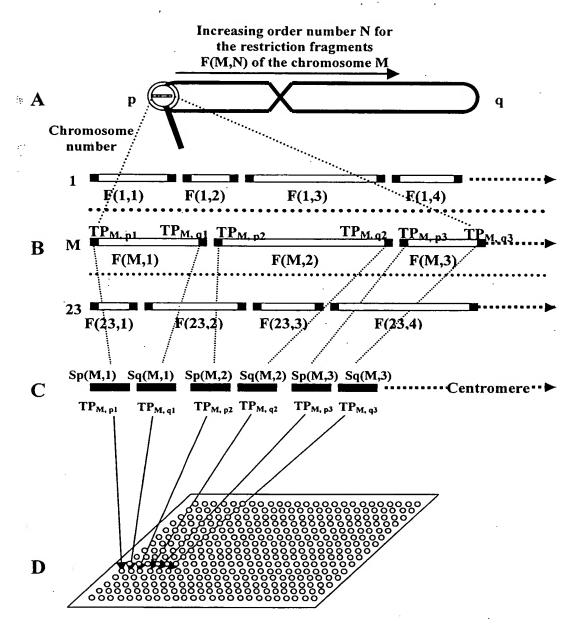


FIG. 36



Micro-array of ordered terminal PENTAmer (TP)
DNA sequences

FIG. 37

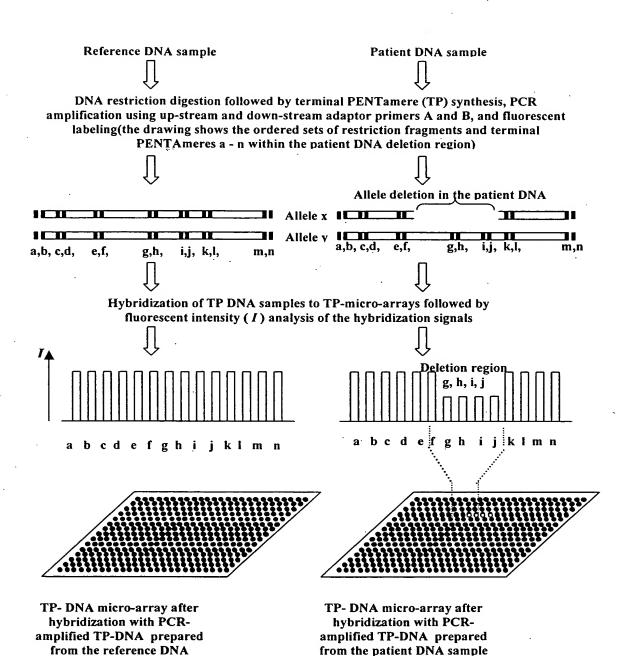


FIG. 38

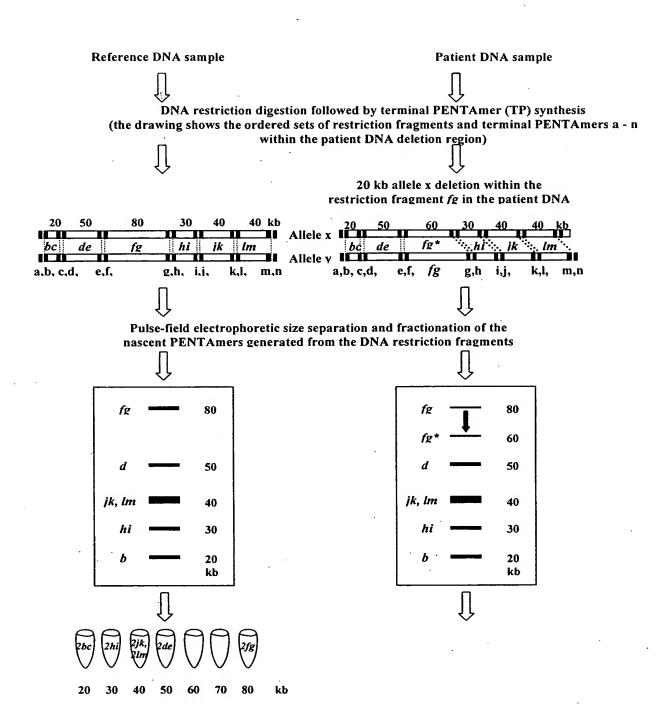
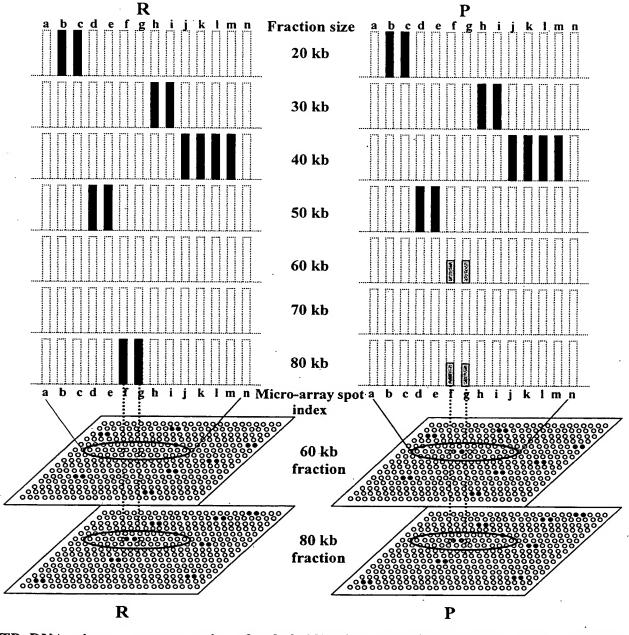


FIG. 39A



Hybridization of TP DNA size fractions to the TP-micro-array followed by fluorescent intensity analysis of the hybridization signals

Fluorescence intensity profiles of the TP- DNA micro-array a-n region after hybridization with PCR-amplified and labeled TP-DNA size fractions 20kb, 30kb, 40kb, 50kb, 60kb, 70kb and 80kb prepared from reference (R) and patient (P) DNA samples



TP-DNA micro-array a-n region after hybridization with PCR-amplified 60kb and 80 kb TP-DNA size fractions prepared from the reference (R) and patient (P) DNA samples

Up-stream, terminus-attaching, nick-translation adaptor A

5' P - GATCGCCTATACCTAGGACCATGTAAddC 3' (SEQ ID NO. 16) 3' ddCGGAUATGGAUCCUGGUACATTG-OH 5' (SEQ ID NO. 17)

Acceptor-adaptor Ac

5' - GATCGCCTATACCTAGGACCATGTAA 3' (SEQ ID NO. 18) 3' CGGAUATGGAUCCUGGUACATTG-OH 5' (SEQ ID NO. 19)

Recombination, nick-translation adaptor RA-(L-cos)

5' P - GATCGCCTATACCTAGGACCATGTAACGAATTCATCA 3' (SEQ ID NO. 20)
NH₂CGGAUATGGAUCCTGGUACATUGCTTAAGTAGTCCCGCCGCTGGA-OH 5' (SEQ ID NO. 21)

Down-stream, nick-attaching adaptors B-3' (a), B-3' (b), B-3' (c) and B-3' (d)

5'-GGGAGATCTGAATTCCCCCCCCCCCCddC-3' (SEQ ID NO. 22)
3'-ddCGCCACTGGGCCCTCTAGACTTAAG - P 5'(SEQ ID NO. 23) (a)

5' -GGGAGATTCTGAATTCAAAAAAAAddA-3' (SEQ ID NO. 26)
3' -ddAGCCACTGGGCCCTCTAGACTTAAG - P 5' (SEQ ID NO. 27) (c)

3- AATGTACCAGGATCCATATCCGCGCCACTGGGCCCTCTAGAC - P 5' (SEQ ID NO. 29) (d)

Oligo-construct with nick

32P-A nick

5'-Biotin-GCGGTGACCCGGGAGATCTGAATTCA GGGCGGCGACCT-3' (SEQ ID NO. 30 & 31)
3'- CGCCACTGGGCCCTCTAGACTTAAGTCCCGCCGCTGGA - P-5' (SEQ ID NO. 32)

a) for a nomenclature of the adaptors A and B-3' see section 6

Fig. 40

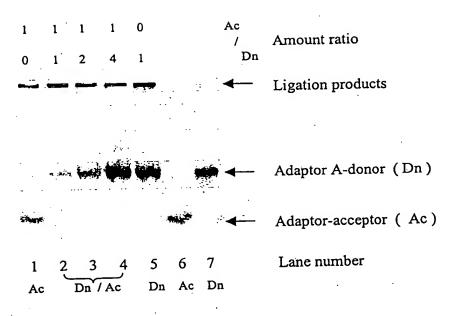


Fig. 41

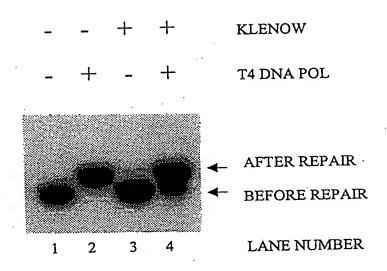
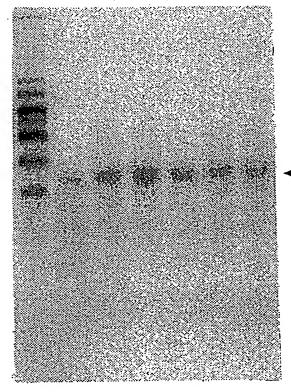


Fig. 42



← 1.4 kb PENT products

M 1 1.5 2 3 5 10

Taq DNA Polymerase, μl

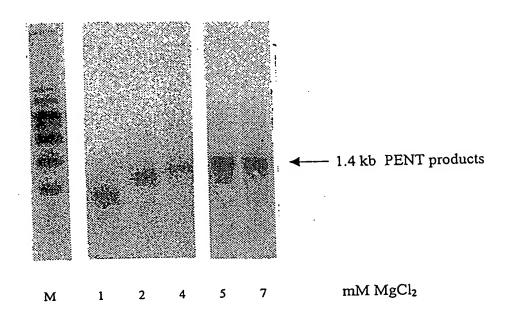
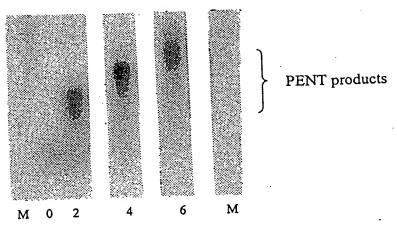


Fig. 44



PENT reaction time, min

Fig. 45

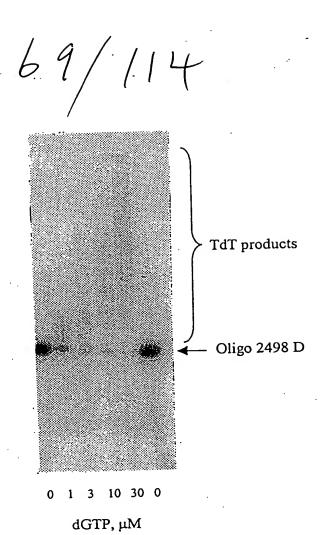


Fig. 46

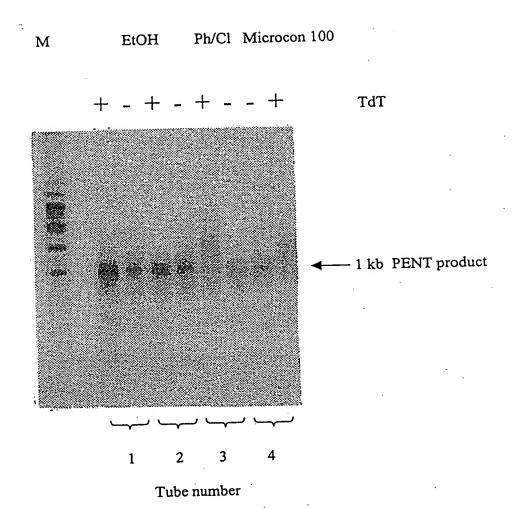


Fig. 47

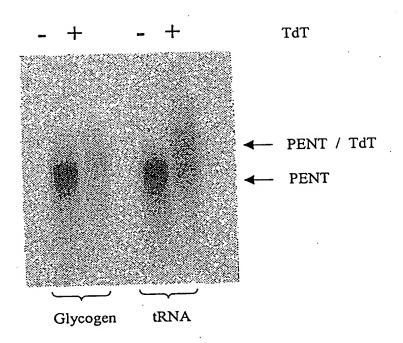


Fig. 48

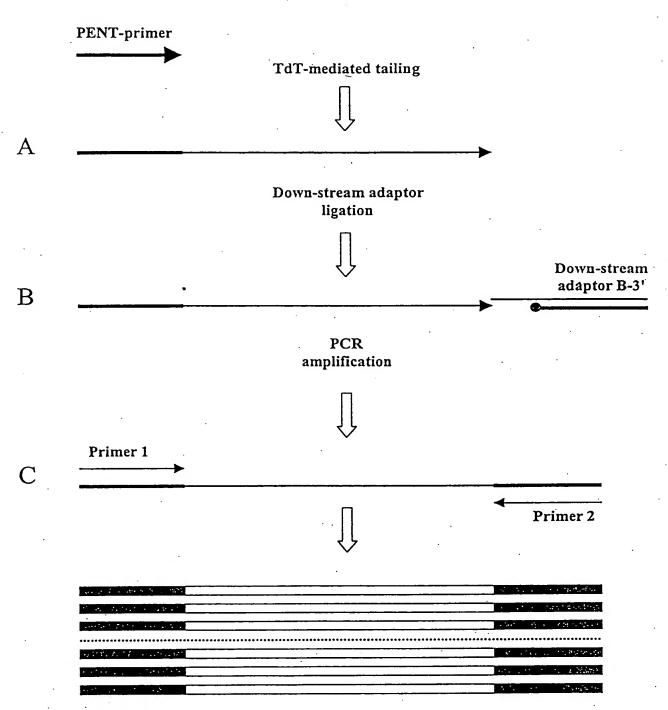
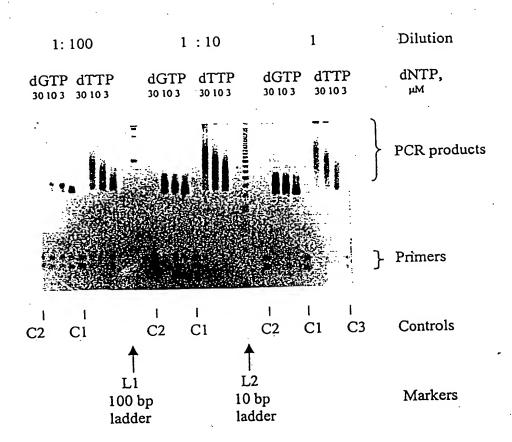


Fig. 49



dTTP dGTP

M 3 10 3 10

dNTP concentration during TdT reaction, μM

← PENTAmer PCR products

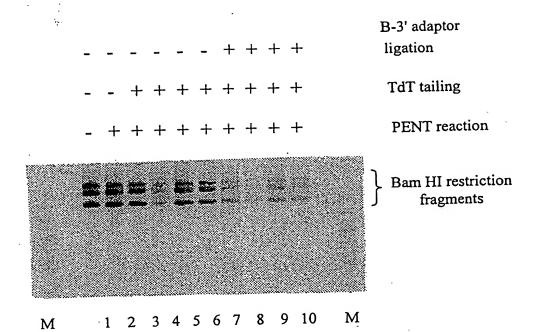


Fig. 52

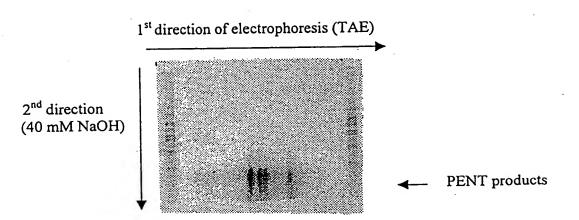
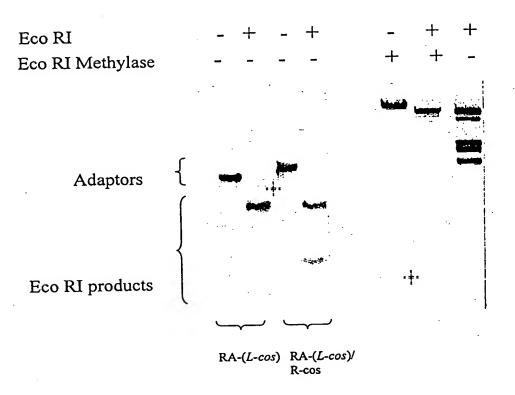


Fig. 53



Oligo-adaptors

Lambda DNA

Fig. 54

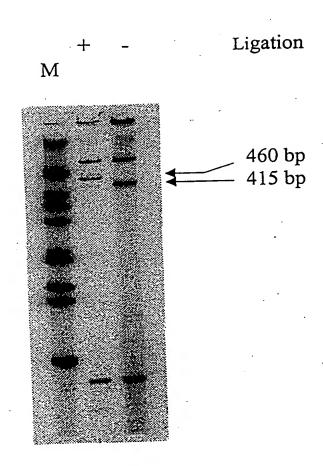


Fig. 55

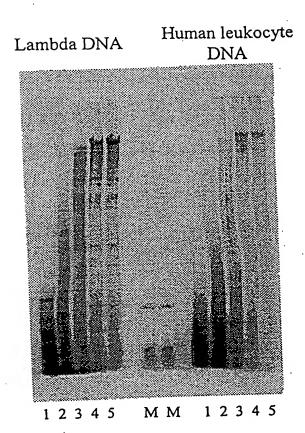
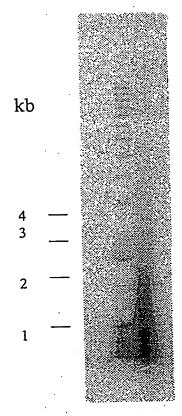
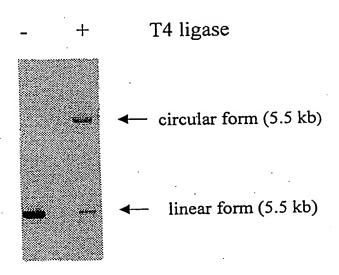


Fig. 56



M DNA

Fig. 57



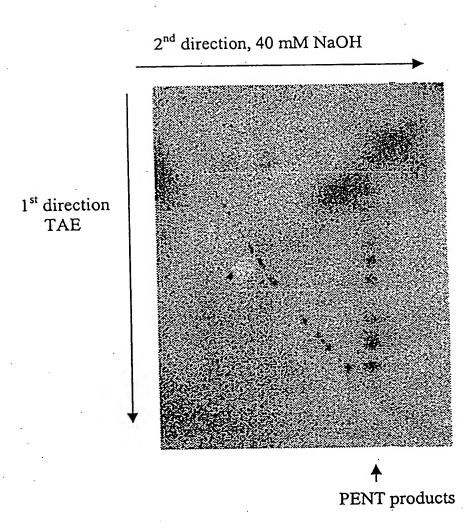


Fig. 59

Lambda DNA methylation protection

 \int

Ligation of the up-stream RA-(L-cos) adaptor to lambda DNA L-cos site

 \iint

Partial digestion with Sau 3A I



DNA circularization by ligation at very low concentration



DNA purification and concentration



Time-controlled PENT reaction initiated at the internal Sau 3A I sites



Recombinant PENTAmer synthesis



DNA linearization with Eco RI



Preparative electrophoretic size fractionation of the nascent PENTAmers



PCR amplification of the PENTAmers



Restriction fingerprint analysis of the amplified DNA fractions

84/114A The RA-(L- $\lambda \cos$) adaptor - Lambda DNA junction structure

 $RA-(L-\lambda \cos)$ adaptor

lambda DNA L-cos end

5' P- GGGCGGCGACCTnnn nnn

5' P- GATCGCCTATACCTAGGACCATGTAACGAATTCATCA -3'OH NH2CGGAUATGGAUCCTGGUACATUGCTTAAGTAGTCCCGCCGCTGGA

B The recombinant junction formed by a circularization reaction.

C The 5'-end of the PENTAmer

nnnnnnnnctagCGGAUATGGAUCCTGGUACATUGCTTAA - 5'

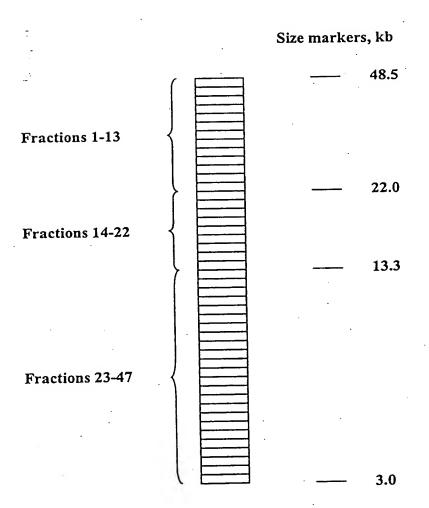
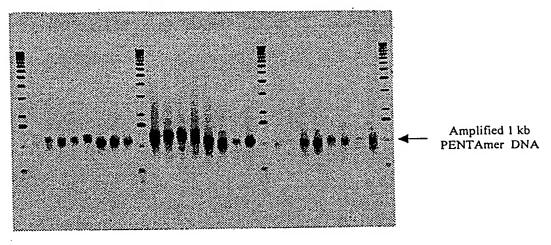


Fig. 62

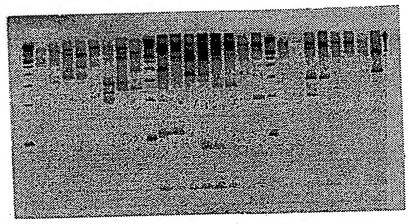


M 1 2 3 4 5 6 7 8 M 9 10 11 12 13 14 15 16 M 17 18 19 20 21 22 23 24 M fraction number

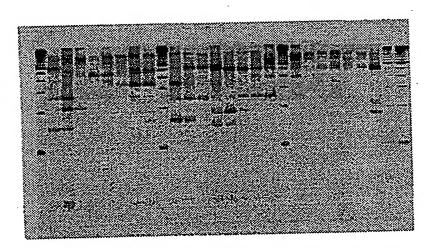
Amplified 1 kb
PENTAmer DNA

M 25 26 27 28 29 30 31 32 M 33 34 35 36 37 38 39 40 M 41 42 43 44 45 46 47 48 M fraction number

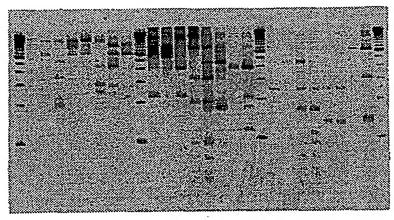
Fig. 63



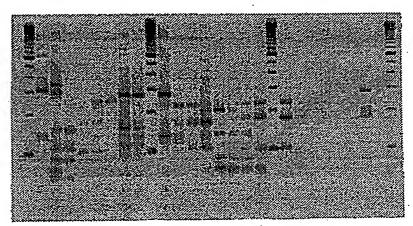
M 1 2 3 4 5 6 7 8 M 9 10 11 12 13 14 15 16 M 17 18 19 20 21 22 23 24 M fraction number



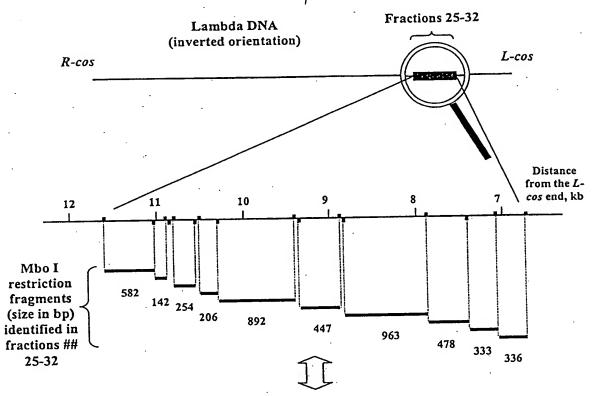
M 25 26 27 28 29 30 31 32 M 33 34 35 36 37 38 39 40 M 41 42 43 44 45 46 47 48 M fraction number



M 1 2 3 4 5 6 7 8 M 9 10 11 12 13 14 15 16 M 17 18 19 20 21 22 23 24 M fraction number



M 25 26 27 28 29 30 31 32 M 33 34 35 36 37 38 39 40 M 41 42 43 44 45 46 47 48 M fraction number



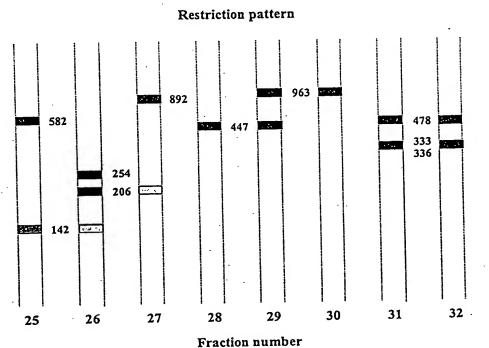
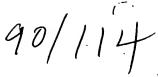


Fig. 66



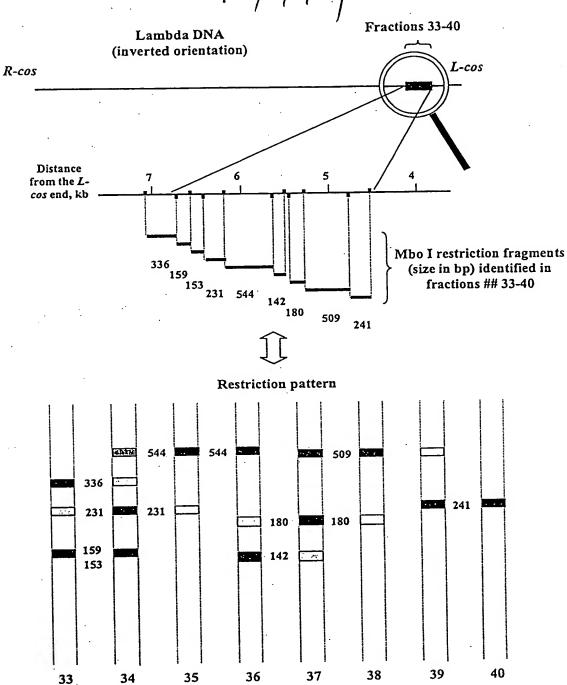


Fig. 67

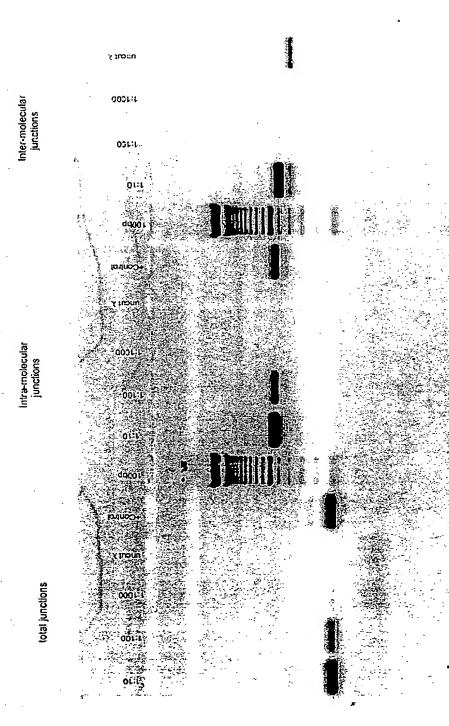
Fraction number

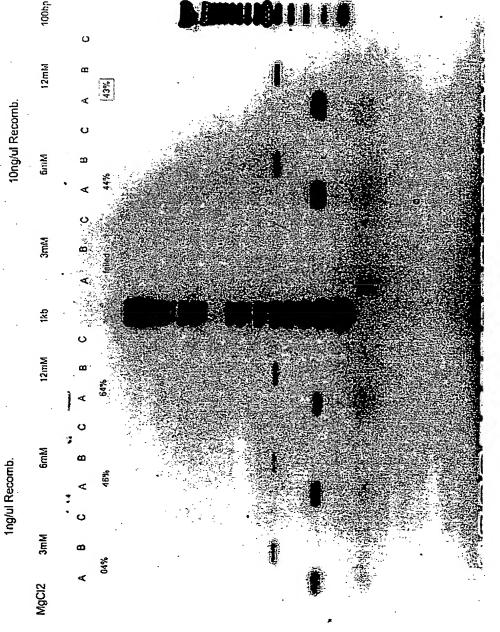


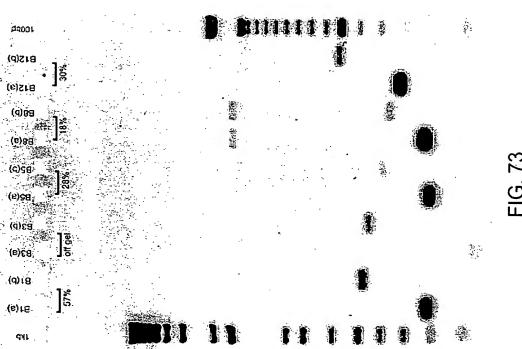
FIG. 68

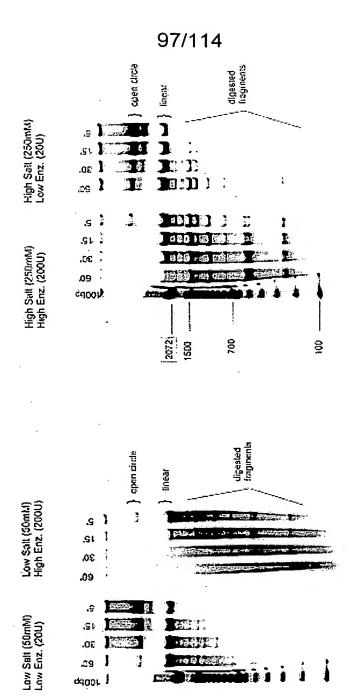
A) RA1/RA2 (Methylation Dpn-I Activation)
RA1 5'-GATCTGAGGTTGTAGAAGACTCGGACGATACATGCACCGTGGTGCAGTCGTAATCCAGTCCCGA (SEQ ID NO:69) AGCCACGTCAGGATTAGGTCAGGGCTCCTGTCGCGATAAAACGATCGGGCACTCGTTATTGATCGCTAG-5' RA2
•
RA2 Component Oligos: RA2(A) 5'-(P)-GATCGCTAGTTATTGCTCACGGGCTAGCAAATAGCGCTGTCCTCGGGACTGGATTACGACTGCACCGATCTC-3' (SEQ ID NO:77) RA2(B) 5'-GAGCAATACTAGC-B3' (SEQ ID NO:78) RA2(C) 5'-(P)-GGACAGCGCTATTTGCTAGCCGGT-3' (SEQ ID NO:79) RA2(C) 5'-(P)-AGAGGGCTTTTGCTAGCCGGTCGTAGCCGGA-B3' (SEQ ID NO:80)
B) Simplified Recombinant Adapters Sral/Sra2
sra 1 5'P-GATCTGAGGTTGTTGAAGACTCGGACGATACACGCTGGGTTGAGGAAGTCGTAAATA TGTGCGACCCAACTCCTTCAGCATTTATTATTGGTAGGGTTGATGGTAG-5'P sra 2
Sra la 5'P-GATCTGAGGTTGTTGAAGACTCGGACGATACACGCTGGGTTGAGGAAGTCGTAAATA-3' (SEQ ID NO:81) Sra lb B3'-ACTCCAACAACTTC-5' (SEQ ID NO:83) Sra lC B3'-ACTCCAACAACTCTGAGCCTGCT-5' (SEQ ID NO:83) Sra lD B3'-TGTGCGACCCAACTCCTTCAGCATTTATTGATGGTTGTCGTTATTGATCGCTAG-5'P Sra 2A (SEQ ID NO:85) 3'-TGTGCGACCCAACTCCTTCAGCATTTATTTATTGTAGGGTTGTCGTAATTGCTAG-5'P Sra 2B (SEQ ID NO:86) 5'-CAGCAATAACTAGC-B3' Sra 2C (SEQ ID NO:87) 5'-ACACGCTGGGTTGAGGAAGTCGTAAATA-B3' Sra 2D (SEQ ID NO:88)
 Sral' Expanded complementarity with original Sra2 P-GATCTGAGGTTGTTGAAGAGTGTTGAGGAAGTGTAAATAAGCATCCCAA P-GATCTGAGGTTGTTGAAGACCCAACTCCTTCAGGATTTATTATTAGTAGGGTTGTCGTTATTGATGATGGTAGGTTGTTATTGATGATGATGGTAG-P5'
5'P-GATCTGAGGTTGAAGACACGCTGGGTTGAGGAAGTCGTAAATAACCATCCCAA-3' Sra 1A' (SEQ ID NO:89) B3'-ACTCCAACAACTTC-5' B3'-TGTGCGACCCAACTCCTTCAGCATTTAT-5' B3'-TATTGGTAGGGTT-5' Sra 1H (SEQ ID NO:90)

Simplified re Sra 1A ID NO:91)	ecombinant adapter (Sra) oligos P5'-GATCTGAGGTTGTTGAAGACTCGGACGATACACACGCTGGGTTGAGGAAGTCGTAAATA-3' (SEQ
Sra 1B	5'-CTTCAACAACCTCA-B3' (SEQ ID NO:92)
Sra 1C Sra 1D	5'-TCGTCCGAGTCTTCAACAACCTCA-B3' (SEQ ID NO:93) 5'-TATTTACGACTTCCTCAACCCAGCGTGT-B3' (SEQ ID NO:94)
Sra 2A NO:95)	P5'-GATCGCTAGTTATTGCTGTTGGGATGGTTATTTATTTACGACTTCCTCAACCCAGCGTGT-3' (SEQ ID
Sra 2B	5'-CAGCAATAACTAGC-B3' (SEQ ID NO:96)
Sra 2C Sra 2D	5'-AACCATCCCAACAGCAATAACTAGC-B3' (SEQ ID NO:97) 5'-ACACGCTGGGTTGAGGAAGTCGTAAATA-B3' (SEQ ID NO:98)
Sra 1A' NO:99)	P5'-GATCTGAGGTTGTTGAAGACACGCTGGGTTGAGGAAGTCGTAAATAAA
Sra 1H	5'-TTGGGATGGTTATT-B3' (SEQ ID NO:100)
	pination screening oligos
Total(+)	5'-AGGTTGTAGAAGACTCGG-3' (SEQ ID NO:101)
Total(-)	5'-GCTAGTTATTGCTCACGG-3' (SEQ ID NO:102)
	5'-GCATCGCTTGAATTGTCC-3' (SEQ ID NO:103)
	5'-TGCTCTCGGAATATCAAT-3' (SEQ ID NO:104)
	5'-GCATCGCTTGAATTGTCC-3' (SEQ ID NO:105)
Inter(-34599)	5'-ATATTCAGGCCAGTTATC-3' (SEQ ID NO:106)
E-coli recomb B1(RP)	vination screening oligos 5'-CTTACACCGGCGAAGTGAAAG-3' (SEQ ID NO:107)
B1 (PCR)	5'-CGCTGCCGGAGCTGTTAGACAATTC-3' (SEQ ID NO:108)
B1 (NEST)	5'-GCCTGCAAGCCGGTGTAGACATCAC-3' (SEQ ID NO:109)
B3(RP)	5'-CTGCAGGCCAGGAGAGAT-3' (SEQ ID NO:110)
B3 (PCR)	5'-GTTGTGGCCTTCCAGTAAGGTCC-3' (SEQ ID NO:111)
B3 (NEST)	5'-GCAAAATAGCTGGCAGGTGTAGG-3' (SEQ ID NO:112)
B5 (RP)	5'-TAGGGCGGCATCAGGTAATAC-3' (SEQ ID NO:113)
B5 (PCR)	5'-TGCCGCCGTTCGCATCCATACCA-3' (SEQ ID NO:114)
B5 (NEST)	5'-TTCCCTGCCTGGTCGCCGTATCTGTG-3' (SEQ ID NO:115)
B8 (RP)	5'-TGAAGGATACGGAAGCAGAAA-3' (SEQ ID NO:116)
B8 (PCR)	5'-GCCATTGCTGATTGCCCACCGACAA-3' (SEQ ID NO:117)
B8 (NEST)	5'-CTCTATCGCTCGGCCTAAGTCTTTAC-3' (SEQ ID NO:118)
B12(RP)	5'-GCGGTCGGCGTGGATAAAGTA-3' (SEQ ID NO:119)
B12 (PCR)	5'-GTGAGCGGGATGAACGAACCTTA-3' (SEQ ID NO:120)
B12 (NEST)	5'-CTGCGCCAGGGCTTCCAGACATTGTG-3' (SEQ ID NO:121)

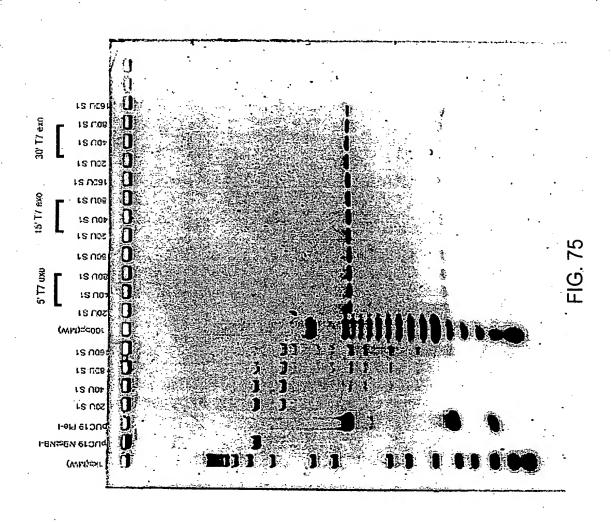


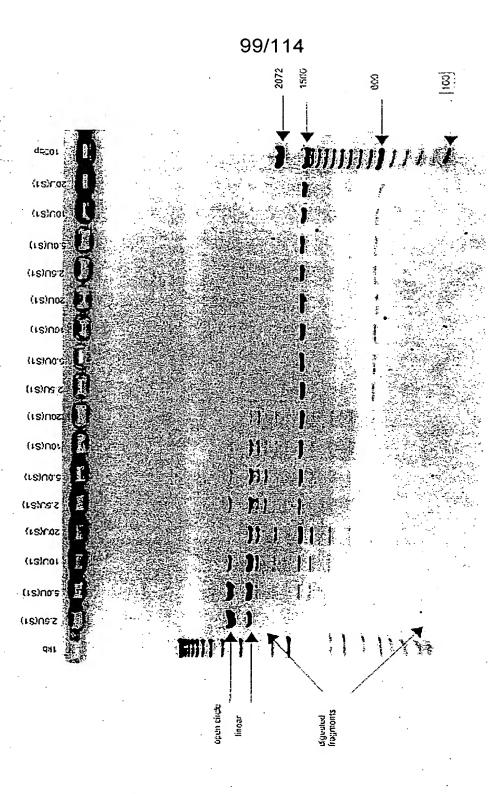


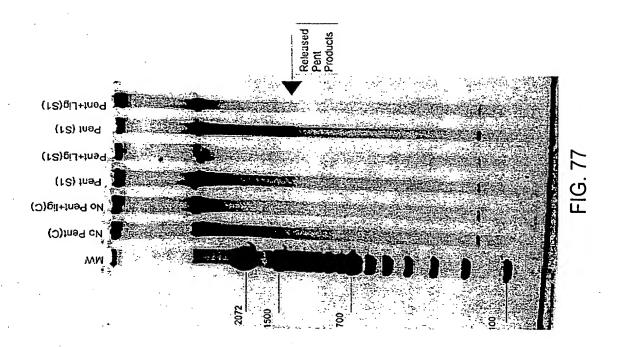


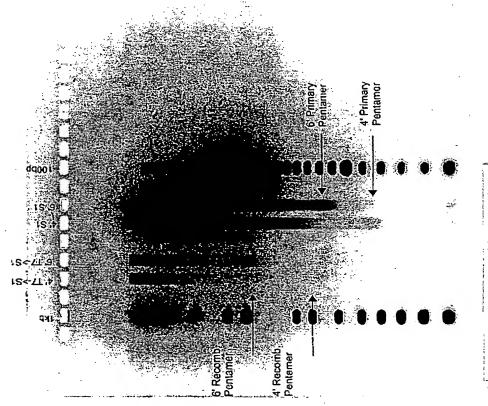


29 400pb

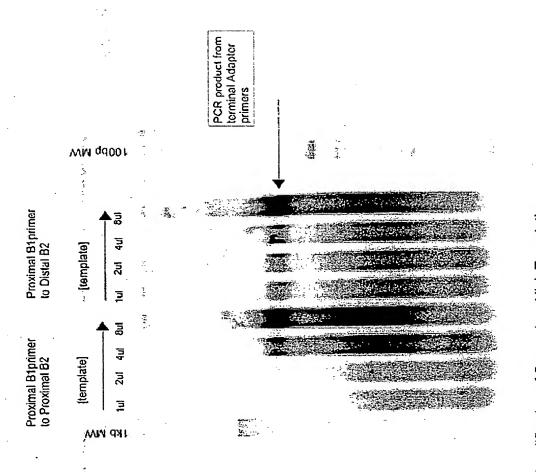




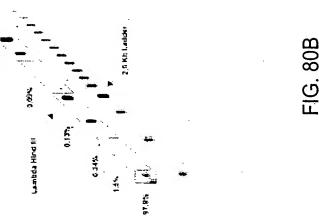


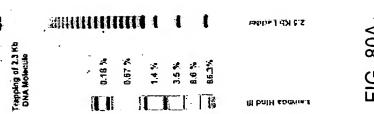


Enzymatic Release of Recombinant Pentamers T7 gene6 - S1 nuclease



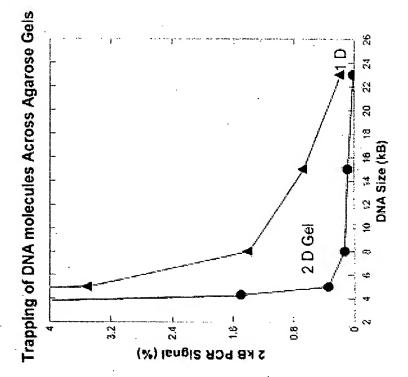
Amplification of Secondary Nick Translation released recombinant Pentamers



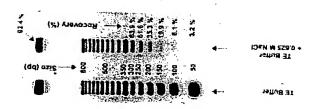


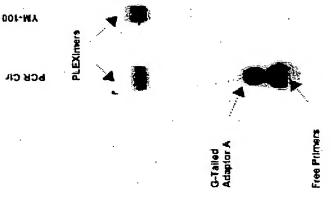
-1G. 80A

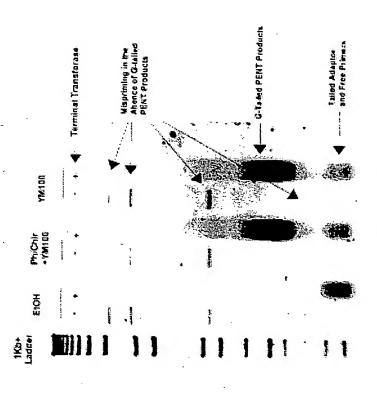


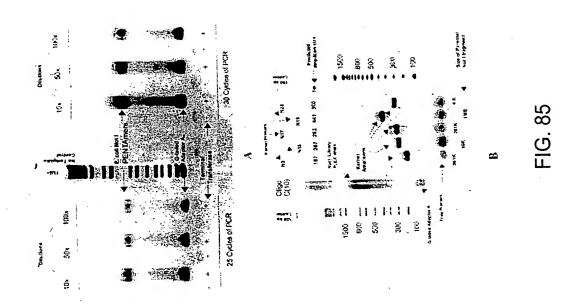


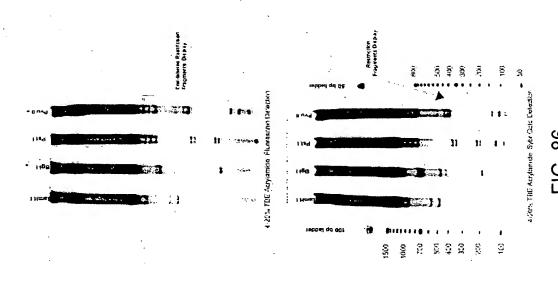
Recovery of DNA Fragments after Microcon YM-100 Filtration

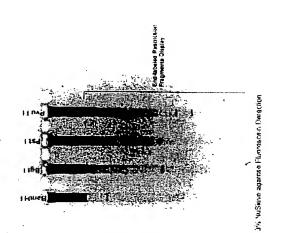


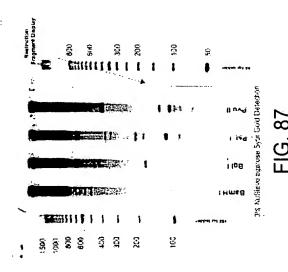








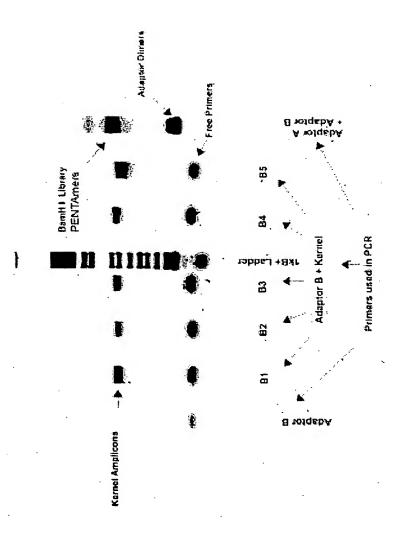


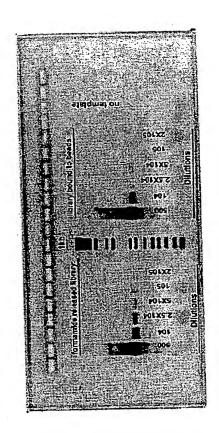


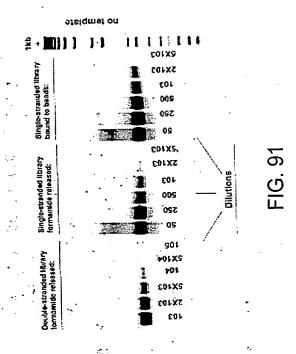
8











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